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## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

<b>(51) International Patent Classification <sup>7</sup> :</b> <b>A61K 39/395, 38/02, C07K 16/18, C12N 1/21, 15/85, C12Q 1/68</b>		<b>A1</b>	<b>(11) International Publication Number:</b> <b>WO 00/10603</b>
			<b>(43) International Publication Date:</b> 2 March 2000 (02.03.00)
<b>(21) International Application Number:</b> PCT/US99/19337			<b>(81) Designated States:</b> CA, JP, European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE).  <b>Published</b> <i>With international search report. Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i>
<b>(22) International Filing Date:</b> 24 August 1999 (24.08.99)			
<b>(30) Priority Data:</b> 60/097,640      24 August 1998 (24.08.98)      US Not furnished      24 August 1999 (24.08.99)      US			
<b>(71) Applicant:</b> THE BOARD OF TRUSTEES OF THE LELAND STANFORD JUNIOR UNIVERSITY [US/US]; Office of Technology Licensing, Suite 350, 900 Welch Road, Palo Alto, CA 94304-1850 (US).			
<b>(72) Inventors:</b> HOPE, Ernest, G.; 1313 Valley Run, Durham, NC 27707 (US). NEGRIN, Robert; 185 Heather Lane, Palo Alto, CA 94303 (US).			
<b>(74) Agents:</b> TRECARTIN, Richard, F. et al.; Flehr Hohbach Test Albritton & Herbert LLP, 4 Embarcadero Center, Suite 3400, San Francisco, CA 94111-4187 (US).			
<b>(54) Title:</b> COMPOSITIONS AND METHODS FOR PROTECTING ORGANS, TISSUE AND CELLS FROM IMMUNE SYSTEM-MEDIATED DAMAGE			
<b>(57) Abstract</b>  This invention concerns novel compositions and methods for the protection of organs and cells from damage caused by activated lymphocytes, NK-cells and NK-like cells, more particularly compositions and methods for the protection of vascular endothelial cells from immune system-mediated damage.			

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**COMPOSITIONS AND METHODS FOR PROTECTING  
ORGANS, TISSUE AND CELLS FROM  
IMMUNE SYSTEM-MEDIATED DAMAGE**

**FIELD OF THE INVENTION**

The field of the invention is compositions and methods for the protection of tissue, organs and cells from immune system mediated damage.

**BACKGROUND OF THE INVENTION**

- 5 Bone marrow transplantation (BMT) represents one form of curative therapy for patients with malignant diseases. The beneficial effects of BMT are not only due to the high dose chemo- and radiation therapy but also to immune mechanisms termed the graft-versus-leukemia effect (GvL) (Sullivan *et al.*, *Blood* 73:1720 (1989); Weiden *et al.*, *N. Engl. J. Med.* 300:1068-1073 (1979)).
- 10 Both natural killer (NK) cells and T-cell subsets contribute to GvL (Antin, *Blood* 82:2273-2277 (1993); Hauch *et al.*, *Blood* 75:2250-2262 (1990)). This is based on the observation that removal of T-lymphocytes from the stem cell product results in dramatically enhanced relapse rates following allogeneic BMT (Martin *et al.*, *Blood* 66:664-672 (1985)). It has also been recently
- 15 demonstrated that infusion of donor lymphocytes into patients, who have relapsed following T-cell depleted allogeneic BM transplant, results in a significant percentage of patients re-entering a complete remission (Collins *et al.*, *Clinical Oncology* 15:433-444 (1997); Kolb *et al.*, *Blood* 86:2041-2050 (1995); Porter *et al.*, *N. Engl. J. Med.* 330:100-106 (1994)). The identification

and *in-vitro* expansion of NK- and T- cell subsets with spontaneous or acquired GvL anti-tumor activity is an area of active investigation.

5 The understanding of biology of classical cytotoxic T lymphocytes (CTL) and their use of TCR  $\alpha/\beta$ , CD3 and CD8 in the context of appropriate MHC I expression on target cells is advanced (Guidos *et al.*, *J. of Exp. Med.* 172:835-845 (1990); Weissman, *Cell* 76:207-218 (1994)). However, the biology of the rarer subsets of cytotoxic T-cells and of NK cells which do not use the above molecules in the sense of classical MHC I restricted cytotoxicity remains a challenge. For example, the field of tumor immunology is  
10 challenged with the consistent finding of unorthodox T-cell cytotoxic activity and expansion in the context of auto and allo reactivity of tumor targets which lacks classical features of MHC I restriction (Hoglund *et al.*, *Immun. Rev.* 155:11-28 (1997)). Several distinct T and NK cell populations have been expanded *in vitro* and utilized in-vivo as effector cells for adoptive immune  
15 therapy.

Lymphokine activated killer (LAK) cells are IL-2 dependent short term culture products with limited proliferative potential derived from NK cells. LAK cells recognize a broad array of tumor cell targets and autologous tumor cells *in-vitro*; however, their *in-vivo* efficacy is limited by the requirement for  
20 the co-application of IL-2 (Blaise *et al.*, *Eur. Cytokine Netw.* 2:121-129 (1991); Fortis *et al.*, *Cancer Immunol. Immunother* 33:128-132 (1991); Lee *et al.*, *J. Biol. Response Mod.* 7:43-53 (1988); Nalesnik *et al.*, *Transplantation* 63:1200-1205 (1997); Rosenberg, *J. Biol. Response Mod.* 3:501-511 (1984); Rosenberg *et al.*, *New England J. of Med.* 316:889-897 (1987); Rosenberg *et al.*, *J. of*  
25 *Exp. Med.* 161:1169-1188 (1985); Teichmann *et al.*, *Leuk. Res.* 16:287-98 (1992). The systemic administration of IL-2 is associated with considerable toxicities which are characterized by endothelial destruction, both *in-vitro* and *in-vivo* (Siegel *et al.*, *J. of Clin. Oncology* 9:694-704 (1991)). LAK cell therapy has led to significant morbidity and even fatalities associated with the vascular  
30 leak syndrome (VLS) (Glauser *et al.*, *Am. J. Med. Sci.* 296:406-412 (1988);

Kotasek *et al.*, *Cancer Res.* 48:5528-5532 (1988); Kozeny *et al.*, *J. Clin. Oncology* 6:1170-1176 (1988); Rosenberg *et al.*, *Ann. Surg.* 210:474-484.

Tumor infiltrating lymphocytes (TIL) require isolation from a surgical specimen. They usually have a high selectivity for their respective tumor.

5 However, their generation tends to be cumbersome and associated with low yields. Specific tumor antigens are rarely available to ensure sufficient *in-vitro* expansion for either successive treatments, or dose escalations (Rosenberg *et al.*, *New. Engl. J. of Med.* 319:1676-1680 (1988); Rosenberg *et al.*, *J. Nat. Canc. Inst.* 86:1159-1166 (1994)).

10 Cytokine Induced Killer (CIK) cells are generated in the absence of target cells from peripheral blood mononuclear cells (PBMC) by *in-vitro* culture in presence of IFN- $\gamma$ , OKT-3, and IL-2. CIK cells have superior *in-vivo* GvL activity as compared to LAK and are capable of purging SCID mice from lethal hematopoietic tumor burden (Lu *et al.* *J. Immunol.* 153:1687-1696  
15 (1994); Schmidt-Wolf *et al.*, *Ann. Hematol.* 74:51-56 (1997); Schmidt-Wolf *et al.*, *Exp. Hematol.* 21:1673-1679 (1993)). Like other forms of activated NK or T cells, bulk cultured CIK show some undesired cytotoxicity against *in-vitro* cultured endothelium (EC).

Vascular leak syndrome (VLS) is a significant side effect associated  
20 with adoptive immunotherapy for treating diseases such as cancer. In particular, it has been reported that the use of IL-2 and the various forms of IL-2 activated NK and T-cells can lead to significant toxicity including VLS. However, no agent has been reported which is capable of inhibiting the lysis of endothelial cells by activated, non-classical MHC class I restricted NK- and  
25 T-cells (Blaheta *et al.*, *Immunology* 94:213-220 (1998); Finnegan *et al.*, *Cancer* 82:186-199 (1998); Utoguchi *et al.*, *Inflammation* 21:223-233 (1997)).

Vascular damage by the immune system is one manifestation of an overall activation of the immune system. While activation of the immune response is extremely important to a host's health and proper functioning, there  
30 are a number of situations where such activation is undesirable. One particular area is associated with transplantation, where one rarely has an identical match

between the donor and recipient of the MHC antigens. Another clinical situation is autoimmune disease. CTLs attack cells where the MHC and associated peptide are both endogenous, as occurs in diseases such as insulin-dependent diabetes mellitus (IDDM). Other undesirable immune system activation events are associated with such maladies as septic shock, rheumatoid arthritis, Crohn's disease, inflammatory bowel disease, asthma, graft versus host disease, coronary artery disease and other cardiomyopathies, adult respiratory distress syndrome, viral liver cirrhosis, and the like.

Immunosuppression has become a general approach in situations where the inhibition of activation of CTLs, NKs and other NK-like cells is desired. However, immunosuppressants such as corticosteroids, cyclosporin A, FK506, and the like, have numerous undesirable side effects including, for example, a general immunosuppressive effect on the host's entire immune system. There is, therefore, substantial interest in identifying new compositions which can specifically protect tissues from damage caused by cells such as T-lymphocytes and other NK-like cells, particularly CTLs, while having less of a universal immunosuppressive effect on the immune system and fewer side effects, so as to leave the host with a substantial proportion of the immune system for protection against adventitious infection. Methods and compositions for achieving such specific immunosuppression are herein provided.

#### **RELEVANT LITERATURE**

Hoppe and Negrin, *Blood*, American Society of Hematology, Thirty-Ninth Annual Meeting, December 5-9 (1997), abstract no. 2019, Jiang *et al.*, *Immunology* 87(3):481-486 (1996); Wada *et al.*, *Jpn J. Cancer Res.* 84(8):906-913 (1993), and Krensky *et al.*, *Exp. Opin. Invest. Drugs* 5:809 (1996).

#### **SUMMARY OF THE INVENTION**

Compositions and methods for protecting cells, organs and other tissues, such as vascular endothelial cells, from damage caused by lymphocytes, NK cells and NK-like cells are herein provided. These compositions find use

either by themselves or in conjunction with other immunosuppressing agents or other therapeutic compounds as a combined therapy. The subject compositions may be contacted *ex vivo* with cells, tissue or an organ. In some embodiments, the then treated material is transplanted into a recipient. The subject  
5 compositions may also be introduced *in vivo* by any convenient means, in sufficient amount to substantially protect cells, organs and/or other tissues from immune system-mediated damage. The subject compositions are preferably Hsp47-related immunoprotective polypeptides or nucleic acids encoding such polypeptides.

10 In a preferred embodiment, cells, tissue or organs are contacted with an immunoprotecting amount of an Hsp47-related immunoprotective polypeptide. In one aspect, the polypeptide comprises the sequence AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AX<sub>4</sub>X<sub>5</sub>X<sub>6</sub>R. In this preferred embodiment, X<sub>1</sub> is V, L, A or T, X<sub>2</sub> is L or H, X<sub>3</sub> is S or V, X<sub>4</sub> is D or E, X<sub>5</sub> is Q, K or R, and X<sub>6</sub> is L or V. In another aspect, the polypeptide  
15 comprises the sequence AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AEQLR, where X<sub>1</sub>, X<sub>2</sub> and X<sub>3</sub> can be any amino acid. In a particularly preferred embodiment the polypeptide comprises the sequence AVLSAEQLR. Polypeptides containing these sequences include the Hsp47 polypeptide disclosed herein as well as contain HLA-A antigens and IL-12 molecules. In a further embodiment, the method comprises contacting  
20 the tissue or organs with an immunoprotecting amount of a composition comprising brefeldin. Examples of the diseased states which cause the immune mediated damage include various autoimmune disease, graft vs. host disease and host vs. graft disease.

The invention also includes chimeric molecules of Hsp47-related  
25 immunoprotective polypeptides and labeled forms thereof. Methods are also disclosed for adoptive immunotherapy wherein an expanded set of T lymphocytes is returned to a patient with an immunoprotecting amount of brefeldin or an Hsp47-related immunoprotective polypeptide.

30 Methods are also disclosed for identifying cells which bind to Hsp47-related immunoprotective polypeptides.

**BRIEF DESCRIPTION OF THE DRAWINGS**

Figure 1 shows the nucleotide sequence and corresponding amino acid sequence of the human Hsp47 cDNA and protein, respectively.

5 Figure 2 discloses related sequences among Hsp47, HLA-A antigens and IL-12 as well as consensus sequences.

Figures 3A, 3B and 3C demonstrate CIK mediated lysis of tumor targets and endothelial cells (EC).

Figure 4 demonstrates CIK tumor lysis and EC lysis as a function of brefeldin A (BFA) concentration.

10 Figures 5A, 5B and 5C show immuno detection of induction of Hsp47 in untreated, 3 $\mu$ g/ml and 10  $\mu$ g/ml BFA treated endothelial cell extracts.

Figures 6A and 6B show FPLC purification of p46 from BFA-induced human umbilical vein endothelial cells (HUVEC) extracts.

15 Figure 7 shows the results from a HUVEC-CIK protection assay with purified, native Hsp47 (p46.5).

Figures 8A and 8B shows the cloning of a huHsp47 gene cassette via PCR directed introduction of cloning sites and electrophoretic separation of expression products.

20 Figure 9 shows that externally provided rHsp47 protects HUVEC cells from CIK-mediated lysis.

Figure 10 shows northern analysis that confirms the upregulation of Hsp47 mRNA by BFA treatment.

Figure 11 shows that recombinant expression of huHsp47 renders HUVEC cells resistant to CIK-mediated lysis.

25 Figure 12 shows that the peptide "AVLSAEQLR" which is shared by both Hsp47 and HLA-A2\*201 protects HUVEC cells from CIK-mediated lysis.

Figure 13 demonstrates graft vs. host disease protection by recombinant Hsp47 in conjunction with bone marrow transplantation.

30 Figures 14A and 14B demonstrate the induction of Hsp47 and other proteins by BFA.



Figures 15A, 15B and 15C depict the structural domains of Hsp47, C-terminal truncation of various domains and the effect thereof on EC cell lysis.

### **DETAILED DESCRIPTION OF THE INVENTION**

5 Methods and composition are provided for protecting cells, organs and tissues, such as vascular endothelial cells, from immune system-mediated damage caused by activated lymphocytes, NK and other NK-like cells. The methods and compositions find use for the treatment of a variety of adverse indications that are mediated by activated immune system cells including, for example, septic shock, scleroderma, rheumatoid arthritis, Crohn's disease, 10 inflammatory bowel disease, colitis, asthma, graft versus host disease, coronary artery disease and other cardio myopathies, adult respiratory distress syndrome, and viral liver cirrhosis, and the like.

The immunoprotecting compositions of the present invention comprise Hsp47-related immunoprotective polypeptides and the drug brefeldin A 15 (1,6,7,8,9,11a,12,13,14,14a-decahydro-1,13-dihydroxy-6-methyl-4H-cyclopent[f]oxacyclotridecin-4-one) (Haerri et al., *Helv. Chim. Acta* 46:1235 (1963), Singleton et al., *Nature* 181 1072 (1958) and Betina et al., *Folia Microbiol.* 7:353 (1962)), a known and commercially available therapeutic.

As used herein, the term "Hsp47-related immunoprotective 20 polypeptides" refers to any polypeptide which has immunoprotecting properties which are similar to that of Hsp47 polypeptide as defined herein. For example, human Hsp47 polypeptide has been shown to protect endothelial cells from lysis from CIK cells. An Hsp47-related immunoprotective polypeptide would be one which shares either qualitatively or quantitatively this or other 25 immunoprotective properties of human Hsp47.

In one aspect of the invention, the Hsp47-related immunoprotective polypeptide additionally is characterized by an amino acid sequence motif which corresponds to the overall consensus sequence as set forth in Figure 1. This can be represented as AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AX<sub>4</sub>X<sub>5</sub>X<sub>6</sub>R. In a preferred embodiment, X<sub>1</sub> 30 is V, L, A or T, X<sub>2</sub> is L or H, X<sub>3</sub> is S or V, X<sub>4</sub> is D or E, X<sub>5</sub> is Q, K or R, and X<sub>6</sub>

is L or V. This can alternatively be represented as

A(v,l,a,t)(l,h)(s,v)A(d,e)(k,q,r)(l,v)R.

5 An Hsp47-related immunoprotective polypeptide can also comprise an IL-12 consensus sequence such as that shown in Figure 2. This can be represented as AX<sub>1</sub>LSAEX<sub>5</sub>X<sub>6</sub>R where X<sub>1</sub> is preferably V, L or T, X<sub>5</sub> is preferably Q, K or R, and X<sub>6</sub> is L or V. This aspect of the invention can also be represented as A(v,l,t)LSAE(q,k,r)(l,v)R.

10 In still a further aspect of the invention, the Hsp47-related immunoprotective polypeptide comprises the consensus sequence as set forth for HLA-A as set forth in Figure 1. This can be represented as AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AEQLR. In this embodiment, X<sub>1</sub> is preferably V or A, X<sub>2</sub> is preferably L or H, X<sub>3</sub> is preferably S or V. This aspect of the invention can also be represented as A(v,a)(l,h)(s,v)AEQLR.

15 In a particularly preferred aspect of the invention, the Hsp47-related polypeptide comprises the consensus sequence for Hsp as set forth in Figure 2. This can be represented as AVLSAX<sub>4</sub>X<sub>5</sub>LR. In this embodiment, X<sub>4</sub> is preferably D or E, and X<sub>5</sub> is preferably K or Q. This aspect of the invention can also be represented as AVLSA(d,e)(k,q)LR.

20 In still another preferred embodiment, the Hsp47-related immunoprotective polypeptide has the sequence AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AEQLR, wherein X<sub>1</sub>, X<sub>2</sub> and X<sub>3</sub> can be any amino acid and Hsp47 polypeptides preferably comprising the sequence AVLSAEQLR.

25 Figure 2 shows the relationship between the peptide sequence spanning residues 96 through 104 of human Hsp47 as compared to similar motifs found in other specific HLA-A antigen molecules and IL-12. Accordingly, in the method of reducing immune mediated damage, any Hsp47-related immunoprotective polypeptide involving HLA-A, Hsp47 and IL-12 and immunoprotecting fragments or variants thereof.

30 As used herein, the term "Hsp47 polypeptide" refers to a polypeptide having the sequence set forth in Figure 1. The term also includes proteins which have at least 70% amino acid sequence identity with the sequence as set

forth in Figure 1, more preferably greater than 90%, most preferably greater than 95% identity with the sequence set forth in Figure 1. The term also refers to a protein encoded by a nucleic acid which is capable of hybridizing with a nucleic acid having the sequence set forth in Figure 1 under moderate or stringent conditions as set forth in more detail hereinafter. The definition of Hsp47 polypeptide also includes immunoprotecting fragments and allelic variations as well as modifications introduced by recombinant technique to substitute, insert or delete one or more amino acid residues.

The entire human Hsp47 cDNA sequence is set forth in Figure 1A. It is within the skill level of the art to prepare purified recombinant or synthetic Hsp47 polypeptides. It is also well within the skill level of the art to identify other Hsp47 cDNA and polypeptides from vertebrates such as mammals and employ those non-human cDNAs and Hsp47 polypeptides in the subject compositions and methods. For example, counterparts to human Hsp47 are known in chicken and rat. As such, Hsp47 polypeptides as used herein include Hsp47 polypeptides and immunoprotecting fragments and variants thereof from all vertebrates, preferably mammals and most preferably humans. With regard to variants and fragments of full-length Hsp47 proteins that find use herein, "immunoprotecting" and grammatical equivalents thereof means that the variant(s) or fragment(s) are capable of reducing or eliminating damage to cells, organs or tissue that is caused by activated lymphocytes, NK cells or NK-like cells as compared to damage to the cells, organs or tissue that would be caused by such cells in the absence of the immunoprotecting agent.

With regard to peptide fragments of the full-length Hsp47 polypeptide or other Hsp47-related immunoprotective polypeptide that will find use herein, those amino acid sequence fragments will generally have at least about 8 consecutive amino acids, usually at least about 10 consecutive amino acids, preferably at least about 15 consecutive amino acids and more preferably at least about 20 consecutive amino acids of the Hsp47 protein sequence. The active sequence may be bonded or non-covalently linked within a chain or as a side chain of other peptides or proteins, for a variety of purposes.

The Hsp47 immunoprotective polypeptides as well as other Hsp47-related immunoprotective polypeptides that find use herein may comprise one or more amino acid substitutions, deletions or insertions. Typical conservative substitutions are shown in Table 1 under the heading of preferred substitutions. If such substitutions do not result in a change in biological activity, then more substantial changes, denominated exemplary substitutions in Table 1, or as further described below in reference to amino acid classes, are introduced and the products screened.

TABLE 1

	Original Residue	Exemplary Substitutions	Preferred Substitutions
	Ala (A)	val; leu; ile	val
	Arg (R)	lys; gln; asn	lys
15	Asn (N)	gln; his; lys; arg	gln
	Asp (D)	glu	glu
	Cys (C)	ser	ser
	Gln (Q)	asn	asn
	Glu (E)	asp	asp
20	Gly (G)	pro; ala	ala
	His (H)	asn; gln; lys; arg	arg
	Ile (I)	leu; val; met; ala; phe; norleucine	leu
	Leu (L)	norleucine; ile; val; met; ala; phe	ile
25	Lys (K)	arg; gln; asn	arg
	Met (M)	leu; phe; ile	leu
	Phe (F)	leu; val; ile; ala; tyr	leu
	Pro (P)	ala	ala
30	Ser (S)	thr	thr
	Thr (T)	ser	ser
	Trp (W)	tyr; phe	tyr
	Tyr (Y)	trp; phe; thr; ser	phe
	Val (V)	ile; leu; met; phe; ala; norleucine	leu
35			

Hsp47 polypeptide-encoding nucleic acids that find use in the subject compositions and methods will have at least about 70%, usually at least about 75%, more usually at least about 80%, preferably at least about 85%, more preferably at least about 90% and most preferably at least about 95% nucleotide sequence identity with the corresponding nucleotide sequence shown in Figure 1A.

"Percent (%) amino acid sequence identity" with respect to the Hsp47 polypeptide sequences identified herein is defined as the percentage of amino acid residues in a candidate sequence that are identical with the amino acid residues in the wild-type human Hsp47 sequence (see Figure 1A), after aligning the sequences and introducing gaps, if necessary, to achieve the maximum percent sequence identity, and not considering any conservative substitutions as part of the sequence identity. "Percent (%) nucleotide sequence identity" with respect to the Hsp47 polypeptide-encoding sequences identified herein is defined as the percentage of nucleotides in a candidate sequence that are identical with the nucleotide sequence shown in Figure 1, after aligning the sequences and introducing gaps, if necessary, to achieve the maximum percent sequence identity. The % identity values used herein are generated by WU-BLAST-2 which was obtained from [Altschul *et al.*, *Methods in Enzymology*, 266:460-480 (1996); <http://blast.wustl.edu/blast/README.html>]. WU-BLAST-2 uses several search parameters, most of which are set to the default values. The adjustable parameters are set with the following values: overlap span = 1, overlap fraction = 0.125, word threshold (T) = 11. The HSP S and HSP S2 parameters are dynamic values and are established by the program itself depending upon the composition of the particular sequence and composition of the particular database against which the sequence of interest is being searched; however, the values may be adjusted to increase sensitivity. A % amino acid sequence identity value is determined by the number of matching identical residues divided by the total number of residues of the "longer" sequence in the aligned region. The "longer" sequence is the one having the

most actual residues in the aligned region (gaps introduced by WU-Blast-2 to maximize the alignment score are ignored).

Hsp47 polypeptides or other Hsp47-related immunoprotective polypeptides that find use herein may be fused to heterologous amino acid sequences in order to provide chimeric or fusion proteins that will find use in the subject invention. Additionally, Hsp47 polypeptides and Hsp47-related polypeptides that find use herein may be obtained from recombinant expression of nucleotide sequences that encode the desired polypeptide(s). In this regard, nucleotide sequences that encode Hsp47 polypeptides of the present invention will hybridize to the nucleotide sequence shown in Figure 1 under moderately stringent conditions, preferably under stringent conditions. As used herein, "stringent conditions" means (1) employing low ionic strength and high temperature for washing, for example, 15 mM sodium chloride/1.5 mM sodium citrate (0.1x SSC)/0.1% sodium dodecyl sulfate at 50°C, or (2) employing during hybridization a denaturing agent, such as formamide, for example, 50% (vol/vol) formamide with 0.1% bovine serum albumin/0.1% Ficoll/0.1% polyvinylpyrrolidone/50 mM sodium phosphate buffer at pH 6.5 with 750 mM sodium chloride, 75 mM sodium citrate (5x SSC) at 42°C. Another example is use of 50% formamide, 5 x SSC (750 mM NaCl, 75 mM sodium citrate), 50 mM sodium phosphate (pH 6/8), 0.1% sodium pyrophosphate, 5 x Denhardt's solution, sonicated salmon sperm DNA (50 µg/ml), 0.1% SDS, and 10% dextran sulfate at 42°C, with washes at 42°C in 0.2 x SSC and 0.1% SDS. Yet another example is hybridization using a buffer of 10% dextran sulfate, 2 x SSC and 50% formamide at 55°C, followed by a high-stringency wash consisting of 0.1 x SSC containing EDTA at 55°C.

"Moderately stringent conditions" are described in Sambrook *et al.*, *supra*, and include the use of a washing solution and hybridization conditions (*e.g.*, temperature, ionic strength, and %SDS) less stringent than described above. An example of moderately stringent conditions is a condition such as overnight incubation at 37°C in a solution comprising: 20% formamide, 5 x SSC (750 mM NaCl, 75 mM sodium citrate), 50 mM sodium phosphate (pH

7.6), 5 x Denhardt's solution, 10% dextran sulfate, and 20 mg/mL denatured sheared salmon sperm DNA, followed by washing the filters in 1 x SSC at about 37-50°C. The skilled artisan will recognize how to adjust the temperature, ionic strength, *etc.*, as necessary to accommodate factors such as probe length and the like.

"Isolated," when used to describe the various Hsp47 polypeptides or Hsp47-related polypeptides disclosed herein, means polypeptide that has been identified and separated and/or recovered from a component of its natural environment. Contaminant components of its natural environment are materials that would typically interfere with diagnostic or therapeutic uses for the polypeptide, and may include enzymes, hormones, and other proteinaceous or non-proteinaceous solutes. In preferred embodiments, the polypeptide will be purified (1) to a degree sufficient to obtain at least 15 residues of N-terminal or internal amino acid sequence by use of a spinning cup sequenator, or (2) to homogeneity by SDS-PAGE under non-reducing or reducing conditions using Coomassie blue or, preferably, silver stain. Isolated polypeptide includes polypeptide *in situ* within recombinant cells, since at least one component of the Hsp47 polypeptide natural environment will not be present. Ordinarily, however, isolated polypeptide will be prepared by at least one purification step. Isolated Hsp47 polypeptides preferably do not contain Hsp47 specific antibodies bound thereto.

An "isolated" nucleic acid encoding Hsp47 polypeptide or Hsp47-related polypeptides is a nucleic acid molecule that is identified and separated from at least one contaminant nucleic acid molecule with which it is ordinarily associated in the natural source of the Hsp47-related polypeptide-encoding nucleic acid. An isolated Hsp47-related polypeptide-encoding nucleic acid molecule is other than in the form or setting in which it is found in nature. Isolated Hsp47-related polypeptide-encoding nucleic acid molecules therefore are distinguished from the Hsp47-related polypeptide-encoding nucleic acid molecule as it exists in natural cells. However, an isolated Hsp47-related polypeptide-encoding nucleic acid molecule includes Hsp47 polypeptide-

encoding nucleic acid molecules contained in cells that ordinarily express Hsp47 polypeptide where, for example, the nucleic acid molecule is in a chromosomal location different from that of natural cells.

The term "control sequences" refers to DNA sequences necessary for the expression of an operably linked coding sequence in a particular host organism. The control sequences that are suitable for prokaryotes, for example, include a promoter, optionally an operator sequence, and a ribosome binding site. Eukaryotic cells are known to utilize promoters, polyadenylation signals, and enhancers.

Nucleic acid is "operably linked" when it is placed into a functional relationship with another nucleic acid sequence. For example, DNA for a presequence or secretory leader is operably linked to DNA for a polypeptide if it is expressed as a preprotein that participates in the secretion of the polypeptide; a promoter or enhancer is operably linked to a coding sequence if it affects the transcription of the sequence; or a ribosome binding site is operably linked to a coding sequence if it is positioned so as to facilitate translation. Generally, "operably linked" means that the DNA sequences being linked are contiguous, and, in the case of a secretory leader, contiguous and in reading phase. However, enhancers do not have to be contiguous. Linking is accomplished by ligation at convenient restriction sites. If such sites do not exist, the synthetic oligonucleotide adaptors or linkers are used in accordance with conventional practice.

The term "antibody" is used in the broadest sense and specifically covers single anti-Hsp47 polypeptide monoclonal antibodies (including agonist, antagonist, and neutralizing antibodies) and anti-Hsp47 antibody compositions with polyepitopic specificity. The antibodies preferably do not bind to the epitopes reactive with MabN6 or Mab SPA470. In one aspect, the antibodies are specific for epitopes defined by the aforementioned consensus sequences. The term "monoclonal antibody" as used herein refers to an antibody obtained from a population of substantially homogeneous antibodies, *i.e.*, the individual



antibodies comprising the population are identical except for possible naturally-occurring mutations that may be present in minor amounts.

The terms "treating", "treatment" and "therapy" as used herein refer to curative therapy, prophylactic therapy, and preventative therapy.

5           The term "mammal" as used herein refers to any animal classified as a mammal, including humans, cows, horses, dogs and cats. In a preferred embodiment of the invention, the mammal is a human.

          Hsp47-related polypeptides or the fragments thereof may be labeled using standard techniques to incorporate radio labels, *e.g.*,  $^{32}\text{P}$ , fluorescent  
10       labels, *etc.* Additional moieties can be added to facilitate purification of the polypeptide either alone or in conjunction with binding to a cell. For example, Hsp47 polypeptides including a fluorescent label facilitates fluorescence activated cell sorting. Additionally, moieties which are capable of interacting with a magnetic field can be linked to Hsp47. In the latter case, magnetic beads  
15       attached to avidin can be used by combining them with biotin labeled Hsp47 polypeptide.

          Hsp47 was identified as a protein induced in endothelial cells treated with brefeldin A. In addition to Hsp47 (identified as the 465 KDa band in Figure 14A and as p47 in Figure 14B), other proteins are induced by brefeldin  
20       A. These include the protein designated p27 in Figure 14B and a faint band above p47 in Figure 14B having a molecular weight of approximately 60-70 KDa. Such additional proteins may also be immunoprotective polypeptides useful in inhibiting the immune response alone or in combination with all the other proteins induced by brefeldin A.

25           The foregoing molecules can be used to identify cells that bind to the Hsp47 polypeptide or fragment thereof. Detecting the presence of such labels can be performed by any conventional technique, such as radiography, fluorescence detection, fluorescence activated cell sorting and divergence in a magnetic field depending upon the particular type of modified Hsp47  
30       polypeptide used.

The subject compositions find use in a variety of ways. For research purposes, they may be used for analyzing the physiological pathway associated with immunoprotection and/or activation and inactivation of T lymphocytes, NK cells and other NK-like immune system cells as well as immunoprotection of various mammalian tissues, including vascular endothelium. For example, one can combine lymphocytes, particularly CTL cell lines having known peptide targets in conjunction with the subject compositions, in the presence and absence of antigen presenting cells to which the CTLs are restricted. After the lysis by the CTLs, one may then separate the activated CTL cells from quiescent CTL cells by means of the marker CD69, which marker is upregulated *in vitro* upon activation. Separation can be achieved using a FACS and a fluorescent labeled anti-CD69.

By isolating the most fluorescent cells, e.g. the highest 25%, one then lyses the cells and isolates proteins associated with the subject markers, e.g. chromatography, non-denaturing electrophoresis, or the like. Alternatively, one separates the proteins using electrophoresis and then uses a Western blot or other technique with the labeled peptides to identify proteins with which the subject peptides bind. Instead of a radiolabel, any other type of label may be employed, normally a small organic molecule, such as biotin, a fluorescer, and the like. Where biotin is used, after separation, avidin may be added, where the avidin is labeled with a label as described previously.

One may also compare lymphocytes which have been combined with antigen presenting cells in the presence and the absence of the subject immunoprotecting compositions. cDNA libraries may be prepared in each instance and representational differential analysis, subtraction, or the like may be employed to detect the differences in expression between the cells which have been activated in the presence and the absence of the subject compositions. One may also determine whether particular subsets of lymphocytes or other immune system cells respond differently from other subsets to the subject immunomodulating compositions by their expression or lack of expression or one or more proteins, particularly surface membrane

proteins. In this way, lymphocytes may be identified which may be removed by leukophoresis or the like, in order to diminish an unwanted lymphocyte attack on tissue.

Depending upon their intended use, particularly for administration to  
5 mammalian hosts, the Hsp47-related immunoprotective polypeptides, *e.g.*,  
Hsp47 polypeptides, may be modified to change their distribution in the blood  
stream, diminish or enhance binding to blood components, enhancing the  
lifetime of the polypeptide in the blood stream, and the like. Immunoprotective  
polypeptides may be bound to these other components by linkers which are  
10 cleavable or non-cleavable in the physiological environment of the blood.  
Immunoprotective polypeptides may be joined at any point of the polypeptide  
where a functional group is present, such as hydroxyl, thiol, carboxyl, amino, or  
the like. Desirably, binding will be at either the N-terminus or the C-terminus.

The Hsp47-related immunoprotective compositions of the present  
15 invention may also be joined to a wide variety of other oligopeptides or  
proteins for a variety of purposes. For example, components of the subject  
compositions may be covalently linked to an immunogen to produce antibodies  
to components of the subject compositions, where the antibodies may serve for  
identification of other peptides having a comparable conformation. In addition,  
20 the antibodies may be used to prepare anti-idiotypic antibodies which may  
compete with the subject proteins or peptides for binding to a target site. These  
anti-idiotypic antibodies may then be used for identifying proteins to which the  
subject proteins and/or peptides bind. Alternatively, the immunoprotective  
polypeptides may be expressed in conjunction with other peptides or proteins,  
25 so as to be a portion of the chain, either internal, or at the N- or C- terminus.  
By providing for expression of the subject proteins and peptides, various post-  
expression modifications may be achieved. For example, by employing the  
appropriate coding sequences, one may provide for lipidation, *e.g.*, prenylation  
or myristoylation. In this situation, the immunoprotective polypeptides will be  
30 bound to a lipid group at a terminus, so as to be able to be bound to a lipid  
membrane, such as a liposome. For administration, liposomes may be used,

where drugs may be introduced into the lumen of the liposome, so as to cooperate with the subject polypeptides in protecting tissues from immune system-mediated damage. Thus, immunosuppressants may be included in the lumen, so that the subject compositions and immunosuppressant may act in a localized manner.

Hsp47-related immunoprotective polypeptides such as Hsp47 may be PEGylated, where the polyethyleneoxy group provides for enhanced lifetime in the blood stream. The Hsp47-derived polypeptides or other immunoprotective polypeptides may also be combined with other proteins, such as the Fc of an IgG isotype, which may be complement binding or not bind complement, or with a toxin, such as ricin, abrin, diphtheria toxin, or the like, particularly the A chain.

The Hsp47-related immunoprotective polypeptides may be modified in a wide variety of ways. Sequence analogs may be prepared by oligopeptide synthesis using a stepwise substitution of the amino acids at each position with alanine or valine, particularly alanine. Generally the total number of amino acids substituted will not exceed 3, ranging from 1 to 3, usually 1 to 2. Methods of producing "scanning" mutations are known in the art, and have been successfully used with a number of different peptides. Examples of protocols for scanning mutations may be found in Gustin, *et al. Biotechniques* 14:22 (1993); Barany, *Gene* 37:111-123 (1985); Colicelli, *et al. Mol Gen Genet* 199:537-539 (1985), and Prentki, *et al. Gene* 29:303-313 (1984).

One can prepare these compositions by preparing a gene coding for an Hsp47-related immunoprotective polypeptide such as Hsp47 polypeptide (see, e.g., Figure 1A). The gene may be introduced into an appropriate expression vector, there being many expression vectors commercially available, whereby the gene is then expressed in an appropriate host. See, Sambrook et al., *Molecular Biology: A Laboratory Manual*, Second Edition, Cold Spring Harbor Laboratories, Cold Spring Harbor, NY, 1989.

The Hsp47-related immunoprotective polypeptides may be prepared by synthesis or by using recombinant techniques, as indicated above. Various

commercial synthetic apparatuses are available, for example automated synthesizers by Applied Biosystems Inc., Foster City, CA, Beckman, etc. By using synthesizers, naturally occurring amino acids may be substituted with unnatural amino acids, particularly D-stereoisomers, side chains having different lengths or functionalities, and the like. For recombinant techniques, one may prepare a nucleic acid sequence which encodes a plurality of the subject polypeptides in tandem, with an intervening amino acid or sequence, which allows for cleavage to the single polypeptide or head to tail dimer. Where methionine is absent, one may have an intervening methionine which allows for single amino acid cleavage. Alternatively, one may introduce consensus sequences, which are recognized by particular proteases for enzymatic cleavage. The particular sequence and the manner of preparation will be determined by convenience, economics, purity required, and the like.

For the most part, the compositions which are used will comprise at least 20% by weight of the desired polypeptide product, more usually at least about 75% by weight, preferably at least about 95% by weight, and for therapeutic purposes, usually at least about 99.5% by weight, in relation to contaminants related to the method of preparation of the product and its purification. Usually, the percentages will be based upon total polypeptide.

By "brefeldin A" is meant the compound having the chemical formula 1,6,7,8,9,11a,12,13,14,14a-decahydro-1,13-dihydroxy-6-methyl-4H-cyclopent[*f*]oxacyclotridecin-4-one, which is commonly known in the art as brefeldin A, as well as any other members of the brefeldin A drug class or other drugs with the same or similar mechanism of action.

By "immunoprotecting amount" is meant the amount of brefeldin A or immunoprotecting polypeptide or expressible nucleic acid encoding it that is capable of reducing or eliminating the lymphocyte-, NK cell- or NK-like cell-mediated destruction of a tissue, preferably a vascular endothelial tissue. Immunoprotecting amounts may differ depending upon the indication for which the composition is employed and may be determined empirically and without undue experimentation by those of ordinary skill in the art.

With regard to brefeldin A, that drug, as well as drugs in the same class and drugs having the same or similar mechanisms of action, are well known and routinely employed in the art (see, e.g., Jiang et al., *supra* and Wada et al., *supra*). Methods and modes for administration of brefeldin A are known in the art and may be employed herein for the protection of tissues from immune system cell-mediated damage.

The subject compositions including brefeldin, Hsp47-related polypeptides and expressible nucleic acids encoding such immunoprotective polypeptides or combinations thereof can be used *in vitro* to inhibit lysis by T or NK cells of target antigen presenting cells, particularly vascular endothelial cells. Thus, in research where one wishes to maintain mixtures of cells, where lymphocytes would be activated and kill antigen presenting cells, such as macrophages or B-lymphocytes, or other cells which might serve as target cells, e.g., neoplastic cells, viral infected cells, or the like, the lysis can be inhibited so that the cellular population may be maintained while under investigation.

The subject compositions including brefeldin, Hsp47-related polypeptides and expressible nucleic acids encoding such immunoprotective polypeptides or combinations thereof can be used may also be used *ex vivo*. In cases of transplantation of organs or cells, more particularly solid organs or particular cells, e.g., bone marrow, whether xenogeneic or allogeneic, the donor organ or cells may be bathed in a medium comprising the subject compositions. In this way, lymphocytes present with the implant will be inhibited from participating in graft versus host disease. Generally, the concentration of the composition will vary in the medium, depending upon the activity of the composition, the level of inhibition desired, the presence of other compounds affecting CTL activation, and the like. Usually, the concentration will be in the range of about 0.1 to 100  $\mu\text{g/ml}$  of polypeptide, more usually in the range of about 1 to 10  $\mu\text{g/ml}$ , although concentrations outside of these ranges may also find use. Concentrations of brefeldin A may be routinely determined empirically, is association with what is known in the art about *in vivo* administration of the drug. Other components of the bathing medium will

generally be constituents normally used in an organ preservation solution, e.g. HBSS. The time for the organ to be maintained in the medium will generally be in the range of about 2 to 72 h.

5 The subject compositions including brefeldin, Hsp47-related polypeptides and expressible nucleic acids encoding such immunoprotective polypeptides or combinations thereof may be also employed *in vivo*,  
administrating the subject compositions by any convenient means for the treatment of autoimmune diseases or to facilitate organ or cellular (*e.g.*, bone marrow) transplants. In the case of transplantation, the subject compositions  
10 may be administered prior to implantation, administration usually beginning not later than about 14 days prior to implantation, there preferably being at least one dosage administered within three days of administration. The subject compositions may be administered in the period beginning about 6 h prior to implantation and may be continued on a predetermined schedule thereafter,  
15 usually not past 30 days, more usually not past 20 days. However, after implantation, the subject compositions may be administered as needed, depending upon the response of the recipient to the organ or cells. In some situations, the subject compositions may be administered chronically, as long as the implant is present in the host. Other forms of *in vivo* use include injection  
20 of the subject compositions into areas of inflammation, *e.g.*, joints, ligaments, tendons and the like as well as various organs such as liver.

Other immunosuppressants which may be present during *in vitro*, *ex vivo* or *in vivo* treatment include injectable, parenteral or topical forms of cortisone, hydrocortisone, modified corticosteroids, Cyclosporin A, FK506,  
25 Imuran, azathioprine, D-penicillamide, MMF (Mofetyl), Methotrexate, cyclophosphamide, gold preparations, salicylates, sulfasalazine, antimalarials and NSAIDs. Biologic agents may also be used in conjunction with the subject compositions, including anti-CD5, Campath Mab, anti-CD4, diphtheria-IL-2 fusion toxin, soluble TNF-R and dimer, soluble IL-1-R, chimeric anti-TNF-  
30 alpha Mab, anti-ICAM-1 Mab, OKT3, anti-thymocyte serum/A6, anti-IL6, antiIL-2, antiCD7, as well as antibodies to CD4, CD8, CD3, LFA-1 and CD28.

Subtherapeutic dosages will be employed, generally when present, not less than about 5% of the normal dosage, and not more than about 75%, usually in the range of about 10 to 60%.

5 In some instances, the subject compositions may be administered in combination with various immunostimulants. Such embodiments exist when the immune system has been compromised, for example, by way of chemotherapeutic agents. In such cases, it is desirable to boost the immune response. The use of the subject compositions in combination with immunostimulants such as IL-2 and other interleukins, interferons, cytokines or  
10 chemokines reduces the side effects associated therewith. Such side effects include the amelioration of vascular leak syndrome which is often associated with treatments with immunostimulants.

Generally, a bolus of the subject composition which is administered will be in the range of about 0.1-50, more usually from about 1-25 mg/kg, of host.  
15 The host may be any mammal including domestic animals, pets, laboratory animals, primates, particularly humans. The amount will generally be adjusted depending upon the half life of the immunoprotecting drug or polypeptide, where the half life will generally be at least one minute, more usually at least about 10 min, desirably in the range of about 10 min to 12 h. Short half-lives  
20 are acceptable, so long as efficacy can be achieved with individual dosages or continuous infusion or repetitive dosages. Dosages in the lower portion of the range and even lower dosages may be employed, where the drug or polypeptide has an enhanced half life or is provided as a depot, such as a slow release composition comprising particles, introduced in a matrix which maintains the  
25 peptide over an extended period of time, e.g., a collagen matrix, use of a pump which continuously infuses the peptide over an extended period of time over a substantially continuous rate, or the like.

The transplantation may involve any organ or cells, including organs such as a heart, kidneys, lung, eyes, liver, gut, vascular vessel, or other organ,  
30 and cells, such as  $\beta$ -islet cells, bone marrow cells, or other cells, where the organ or cells are allogeneic or xenogeneic, particularly where one or more of



the Class I or II MHC antigens are different in the donor as compared to the recipient.

The subject immunoprotecting compositions including brefeldin, Hsp47-related immunoprotective polypeptides and nucleic acids encoding them, by themselves as conjugates or as combinations thereof, may be prepared as formulations in pharmaceutically acceptable media, for example, saline, PBS, aqueous ethanol, glucose, propylene glycol, or the like or as solid formulations in appropriate excipients, generally at a pharmacologically effective dose. The concentrations of the brefeldin A, Hsp47 polypeptide or other immunoprotective polypeptide will be determined empirically in accordance with conventional procedures for the particular purpose. The formulations may include bactericidal agents, stabilizers, buffers, or the like. The amount administered to the host will vary depending upon what is being administered, the purpose of the administration, such as prophylaxis or therapy, the state of the host, the manner of administration, the number of administrations and the interval between administrations, and the like. In order to enhance the half life of the subject compositions, the compositions may be encapsulated, introduced into the lumen of liposomes, prepared as a colloid, or other conventional technique may be employed, which provides an extended life time of the compositions *ex vivo* or *in vivo*. In general, doses of immunoprotective amounts can be packaged in liquid or solid (*e.g.*, lyophilized) form in appropriate containers such as vials, *etc.* If liquid, the composition is preferably in a pharmaceutically acceptable medium (carrier). If solid, it should be prepared so that when reformulated as a liquid (*e.g.*, with water or a pharmaceutical carrier) a pharmaceutically acceptable composition is formed.

The examples disclose the effect of brefeldin A and Hsp47 polypeptides including the polypeptide AVLSEQLR on CIK cell lysis of cultures of endothelial cells derived from human umbilical cord samples. These experiments demonstrated that CIK lysis of the endothelial cells is inhibited by brefeldin A and by at least Hsp47 which is expressed upon

contacting the endothelial cells with brefeldin A. The following discussion summarizes the examples set forth herein as well as other experimental results. However, such results are merely exemplary of the scope of the invention in that the methods and compositions of the invention can be used to meliorate not only the undesirable lysis of endothelial cells but immune mediated damage to other cells, tissues and organs. Moreover, the effect of such compositions and methods is not limited to CIK cells but rather to non-MHC I restricted cytotoxic T lymphocytes and natural killer cells in general. In this regard, it should also be noted that while the methods and compositions of the invention inhibit non-MHC I restricted CTLs, other modes of immune response are not significantly affected. In this regard, a BMT experiment was performed on full body irradiated mice treated with or without an Hsp47 polypeptide. Graft vs. host disease appeared in the non-Hsp47 treated group. Of the members of the Hsp47 treated group, none developed graft vs. host disease. Moreover, none developed opportunistic infection over the time period tested which otherwise would have been expected if the entire immune system was suppressed.

The examples demonstrate that allogeneic and autologous CIK have a high cytotoxicity against a variety of hematopoietic and solid tumor targets including lymphoid, myeloid, and solid tumors as well as against *in-vitro* cultured EC. CIK also lyse additional cancer targets (Lopez *et al.*, *Faseb. J.* 9:A1024 (1995); Lu *et al.*, *J. Immunol.* 153:1687-1696 (1994); Mehta *et al.*, *Blood* 86:3493-3499 (1995); Schmidt-Wolf *et al.*, *Ann. Hematol.* 74:51-56 (1997)). All of the targets studied express normal levels of MHC I.

Anti-MHC class I Mab W6/32, an antibody with proven capability to prevent or disrupt MHC I/TcR interactions *in-vitro* (Shields *et al.*, *Tissue Antigens* 51:567-570 (1998)) does not block CIK mediated target lysis. The nature of the CIK mediated GvL effect was demonstrated to be a purely autologous process, in <sup>51</sup>Cr release assays using cord-blood derived CIK and human umbilical cord derived endothelial targets from the same donor. This is consistent with the concept that CIK represent non classical MHC I restricted

cytotoxic T-cells and excludes the possibility that this observed lysis represents the result of allo-recognition.

CIK are remarkable for their strong GvL effect without causing measurable GvHD in murine models, a finding also confirmed in man. This finding was documented with the SCID/hu *in-vivo* tumor purging model for B-cell lymphoma SU-DHL4 and chronic myelogenous leukemia (Lu *et al.*, *J. Immunol.* 153:1687-1696 (1994); Hoyle *et al.*, *Blood* 92:3318-3327 (1998)). In the examples, a murine SCID system with orthotopic full-thickness human skin allografts was used to provide human cell adhesion molecules in the vessel walls of the skin grafts. This demonstrated lack of allorecognition and GvHD potential by CIK. The established skin grafts did not show signs of CIK infiltration, inflammation or GvHD. Only the human derived neoangiogenesis of solid tumors becomes the selective target of CIK mediated immune surveillance, but not physiologic normal vascular beds outside of the tumor vicinity.

Distinct differences in CIK target recognition become apparent in the analysis of cell adhesion molecules. Our studies demonstrate that the interaction of ICAM-1 on tumor targets with LFA-1 on the CIK effector cells is crucial for the process of CIK mediated lysis of tumor targets. This ICAM-1 / LFA-1 interaction appears not to be involved in CIK mediated EC lysis. These findings are consistent with previous reports on CD3<sup>+</sup>56<sup>+</sup> T-lymphocyte requirements for target cell adhesion (Lu *et al.*, *J. Immunol.* 153:1687-1696 (1994); Schmidt-Wolf, *Cellular Immunology* 169:85-90 (1996)) as opposed to their interaction with EC. CTLs utilize ICAM-1 and LFA-1 in their interaction with tumor targets. NK cells in the absence of MHC I/TcR ligation lyse targets which do not provide sufficient non-kill signalling, in situations where ICAM-1/LFA-1 provides the only identified stimulatory signal. The mechanism of T-cell co-stimulation through ICAM-1/LFA-1 ligation was recently discovered to involve myosin motor protein mediated accumulation of co-stimulatory signalling molecules in the T-cell/tumor target interface, leading to an amplified and prolonged signalling process (Wulfing *et al.*, *Science*

282:2266-2269 (1998)). This process can be captured in real time via syn-capping micro-videography (Wulfing *et al.*, *Proc. Natl. Acad. Sci USA* 95:6302-6307 (1998)). This ICAM-1/LFA-1 interaction is also crucial for the process of lymph node homing at high endothelial venules (Lawrence *et al.*,  
5 *Eur. J. Immunol.* 25:1025-1031 (1995); Oppenheimer-Marks *et al.*, *J. of Immunology* 145:140-148 (1990); Rosenman *et al.*, *J. of Leukocyte Biol.* 53:1-10 (1993)). The ICAM-1/LFA-1 interaction seems to be not required for their interaction with non-activated EC, as this process was found to lack capping of LFA-1 and TcR in their interaction with EC (Kozeny *et al.*, *J. Clin. Oncol.*  
10 6:1170-1176 (1988)). The presence of a ICAM-1/LFA-1 dependent tumor lysis and an ICAM1/LFA-1 independent EC lysis process suggests the existence of at least two distinct pathways of CIK mediated cytotoxicity.

Single cell cloning of CIK demonstrated for the first time that endothelial lysis of activated non-classical MHC I restricted killer T-cells with  
15 anti-tumor activity is a clonal property of the activated T-cells. The mechanisms of cytotoxic endothelial and tumor recognition segregate independently from each other with individual CD3<sup>+</sup>56<sup>+</sup> CIK clones expressing none, either one of the two, or both target recognition capacities. The vast majority of the CIK clones are  $\alpha/\beta$  T-cells, but we also demonstrated that  $\gamma/\delta$   
20 T-cells can differentiate into cytotoxic CIK.

Lack of lytic ability by CIK of either the tumor or the EC target can be the result of at least three different clonal constellations: Each target class may require a specific cascade of signalling molecules to recognize it; lack of certain signalling cascade members in a particular CIK clone may lead to  
25 ineffective signalling for a given target. There may alternatively be a number of independently inhibitory signalling pathways interfering with the execution of a cytotoxic recognition event; protection from CIK lysis may result from a target cell's ability to express a respective ligand. Finally, sensitivity to a particular CIK killing mechanism may differ between tumor targets and EC;  
30 therefore individual clonal expression patterns and levels of alternate target attack mechanisms may account for the observed difference in effective target

lysis. FACS analysis shows that targets express high levels of MHC I, and that functional CIK express  $\alpha/\beta$  TcR (>95%) or  $\gamma/\delta$  TcR (<5%), CD3/CD5 (>98%), CD8 (>80%) or CD4 (<15%), KIR molecules (5-8% staining for DX9) and CD56 (>25% in bulk cultures, >98% of functional clones) (Lu *et al.*, *J.*

5 *Immunol.* 153:1687-1696 (1994)). However, we have not been able to demonstrate functional involvement in target recognition and lysis by CIK for any cell adhesion molecules other than the above mentioned LFA-1 and ICAM-1 nor for the major T- and NK-surface proteins summarized in Table 2.

Effector cells were treated with CMA and BFA (brefeldin A) (Kataoka  
10 *et al.*, *J. Immunol.* 156:3678-3686 (1996)) prior to  $^{51}\text{Cr}$  release cytotoxicity assay to estimate the contribution of perforin/granzyme versus Fas-based CIK cytotoxicity to these targets. Our results demonstrate that EC lysis is virtually entirely perforin and granzyme based, whereas tumor targets are killed utilizing both pathways. Previous research has demonstrated that CIK mediated lysis of  
15 tumor targets is cAMP and  $\text{Ca}^{2+}$  dependent (Mehta *et al.*, *Blood* 86:3493-3499 (1995)).

The combined results from the FasL surface expression inhibition study with BFA and the perforin depletion study with Concanamycin A indicate that cytotoxicity against tumor targets is mediated via granzyme/perforin with  
20 significant contribution of the Fas/FasL pathway. EC lysis, in contrast, is mediated exclusively via granzyme/perforin. This result confirms that differences exist between EC and tumor target physiology which contribute to its sensitivity to CIK.

It is possible that CIK are anergic limited auto-reactive T-cells  
25 (Schwartz, *Curr. Opin. Immunol.* 9:351-357 (1997)). If so, they would represent a pool of T-cells with a heterogenic, differentiated CD complement, which may or may not have lost their function at that time point (Guidos *et al.*, *J. of Exp. Med.* 172:835-845 (1990)). CIK culture like conditions might lead to rescue of anergic T-cells by providing high dose IL-2 stimulation and thus  
30 overcoming anergy (Madrenas *et al.*, *Proc. Natl. Acad. Sci. USA* 93:9736-9741 (1996)).

Given the plurality of surface markers expressed on equally functional clones, the existence of a specific lineage of CIK T-cells and preferential expansion from a distinct precursor cell type found in peripheral blood appears unlikely. Rather, the presence of surface markers on T- and NK cells which lost their effective signalling capacity when tested functionally, has been reported as a hallmark characteristic of anergic cells (Fink *et al.*, *J. Immunol.* 152:4270-4281 (1994)). Our understanding of the biology of T- and NK cells in the anergic state is still limited. But it has been implicated that high levels of IL-2 allow previously anergic T- and NK cells to resume cell division and to develop new, no longer auto-reactive, functional specificities (Alters *et al.*, *Transplantation* 56:633-638 (1993); Hoglund *et al.*, *Immunological Reviews* 155:11-28 (1997); Karpus *et al.*, *Int. Immunol.* 6:721-730 (1994); Karre, *Semin. Cancer Biol.* 2:295-309 (1991); Ljunggren *et al.*, *Immunol. Today* 11:237-244 (1990); Madrenas *et al.*, *Proc. Natl. Acad. Sci. USA* 93:9736-9741 (1996); Nossal, *Ann. N. Y. Acad. Sci.* 690:34-41 (1993); Rothenberg, *Adv. Immunol.* 51:85-214 (1992)). The high doses of IL-2 provided in CIK culture may thus rescue a heterogeneous pool of anergic PBMC T-cells which in the rescue process do not regain their prior potential for auto-reactivity. In addition to the de-novo expression of effector molecules involved in newly gained antitumor functions, CIK may therefore simply maintain their previous individual cell surface expression patterns of now non-functional receptors. This model allows for the observed heterogeneity of FACS markers present on equally functional CIK, as well as the loss of function associated with TcR/MHC I, CD4 and CD8 observed in our functional analysis.

Further insight into the process of cytotoxic T- and NK interaction with EC came from the finding that BFA specifically blocked EC lysis in a dose responsive fashion. This is the first report of an agent capable to suppress EC lysis by activated NK or T-cells. BFA induces a resistant EC state which persists for at least six hours after removal of the agent.

BFA treatment has been reported to lead to a complex series of changes in treated cells, via selective intoxication of a G-protein responsible for protein

transport between cis- and medial golgi compartments. This transport block results in a backing-up of secretion directed proteins into an enlarging ER system and leads finally to the fusion of the cis-golgi compartments with the ER (Lippincott-Schwartz *et al.*, *Cell* 56:801-813 (1989)). The enlargement of the ER compartment increases the synthesis of ER resident proteins. ER resident proteins are characterized by a C-terminal four amino acid "KDEL" or "RDEL"-ER retention signal. Examples of such ER-resident proteins are Hsp47, calreticulin, Grp78 and Grp94 (Ferreira *et al.*, *Arch. Virol.* 138:273-285 (1994); Ferreira *et al.*, *J. Cell Biochem.* 56:518-526 (1994); Ferreira *et al.*, *Connect Tissue Res.* 33:265-273 (1996); Smith *et al.*, *J. Biol. Chem.* 270:18323-18328 (1995)). These ER-resident proteins are not secreted out of the ER compartment, because they are in a  $\text{Ca}^{2+}$  dependent fashion recognized, bound to and re-cycled back into the ER by the KDEL/RDEL-receptors, Erd2.1 and Erd2.2. The ER serves as major intracellular  $\text{Ca}^{2+}$  store. Enlargement of the ER by BFA treatment perturbs the cellular  $\text{Ca}^{2+}$  homeostasis and leads to inactivation of the KDEL/RDEL receptor (Llewellyn *et al.*, *Biochemical and Biophysical Research Communications* 240:36-40 (1997)). ER resident proteins, including Hsp47, thus become freely secreted (Hu *et al.*, *J. Cell Biochem.* 59:350-367 (1995); Hu *et al.*, *J. Cell Biochem.* 59:214-234 (1995)). BFA treatment of EC therefore leads to an increased production of Hsp47 and to the free secretion of this otherwise ER resident protein. Under physiologic conditions, the transport of Hsp47 to the outside of the cell is linked to the expression and co-transport of chaperoned collagens I and IV (Yamamura *et al.*, *Biochem Biophys. Res. Comm.* 244:68-74 (1998)).

$^{35}\text{S}^*$  pulse-chase labelled protein extracts of the BFA induced resistant state of EC were compared with the untreated sensitive phenotype. Amongst the consistently radio-labelled proteins increased in extracts from BFA treated EC was a protein of 46.5 KDa, p46.5. Using phase partitioning at 0-4°C with Triton-X-114 we were able to demonstrate p46.5 to be at least partially localized to the cell membrane of treated EC. Biotinylation of the surface proteins of intact BFA treated EC monolayers with the water soluble, lipid

insoluble reagent biotin-SS-NHS also demonstrated Hsp47 and a 25-27 KDa associated protein to be upregulated at the cell surface after BFA treatment. See Figures 14A and 14B. Biochemically, p46.5 bound to and could be purified with gelatin-sepharose, eluting in both, competitive RGD peptide binding and a low pH elution buffers during FPLC preparations. All biochemical characteristics of p46.5 matched data published for rodent Hsp47 (Hirayoshi *et al.*, *Mol. Cell Biol.* 11:4036-4044 (1991); Jain *et al.*, *Biochem J.* 304:61-68 (1994); Nandan *et al.*, *Biochem Cell Biol.* 68:1057-1061 (1990); Vaillancourt *et al.*, *Biochem J.* 274:793-798 (1991)). Functional analysis demonstrated that gelatin purified p46.5 shares with BFA treatment the unique ability to protect previously CIK sensitive EC in a dose responsive fashion. This was shown through pre-incubation of EC with p46.5 prior to <sup>51</sup>Cr cytotoxicity assay. This is the first report of an externally added protein agent which protects EC from non-MHC I restricted cytotoxicity. This p46.5 protein cross-reacted in immuno-precipitations with antisera raised against membrane proteins from the resistant state of lymphoblastoid cells, and with purified pan-anti heat shock protein monoclonal antibody N6.

p46.5 was positively identified to be human Hsp47 on the original membrane of the N6 immunoprecipitation via the specific anti-Hsp47 monoclonal antibody SPA-470.

The full sequence of human *hsp47* was first described by Clarke and Sandwal after PCR based isolation of partial, but overlapping, cDNA gene fragments spanning the entire human *hsp47* gene (Clarke *et al.*, *Biochem Biophys Acta* 1129:246-248 (1992)). Two plasmids were provided by Dr. Sanwal, corresponding to overlapping partial *hsp47* cDNA fragments of 350 and 1300 bp, respectively. The smaller fragment also contained 5'untranslated regions not coding for the mature gene product of the *hsp47* gene. Within the area of overlap between the fragments there is no shared restriction enzyme sites. PCR based site directed mutagenesis was used to both amplify and ultimately join the two cDNA fragments into one continuous reading frame coding for the human Hsp47 protein. Based on the published sequence data



and our choice of cloning sites, two sets of primers were designed for PCR. An artificial BamHI site was introduced flanking the 5' end of the coding sequence. An artificial EcoRI site was introduced at the 3' end of the coding sequence, as well as a MscI site within each side of the area of overlap between the two clones to facilitate the joining of the cDNAs. After PCR amplification of the two plasmids, the smaller fragment was MscI restricted and gel purified. This smaller fragment was used as the 5' primer on the 1300 bp fragment as template and thus generated the first reported expressible gene cassette of human *hsp47*. The PCR product of this reaction was then directionally ligated into BamHI/EcoRI restricted bacterial protein expression vector pGEX-4T1, and *in-vivo* amplified from single colonies of transformed BL21(DE3) bacterial host cells.

Human *huhsp47* was cloned without its signal peptide "MRSLLLGTLCLLAVALA". pGEX-4T bacterial protein expression system was used for IPTG induction to overexpress Hsp47 and to obtain a fusion protein of human Hsp47 and N-terminally tagged glutathione-S-transferase (GST). The latter allowed use of a one-step affinity purification protocol on GST substrate columns with crosslinked reduced glutathione and elution through competitive binding of this fusion partner with free, soluble reduced glutathione. GST itself has been shown to be non-toxic in most cellular assays and did not increase or reduce target cytotoxicity in the <sup>51</sup>Cr release assays. Purified recombinant huHsp47 protects EC in a dose responsive fashion.

A Kozak initiation sequence and a functional secretion signal for optimal eukaryotic expression was introduced by a combination of PCR and restriction enzyme cloning, using the eukaryotic baculovirus protein expression vector pMel-Bac as template for the Kozak and the mellitin secretion leader sequences. Selection of the strong CMV<sup>IE</sup> promoter was done to ensure a high level of protein expression. The BFA treatment of EC resulted in a marked increase of secreted, extracellular Hsp47, despite the presence of an ER-retention signal "RDEL" on Hsp47. As detailed above, BFA treatment leads to Ca<sup>2+</sup> flux perturbation of the ER compartment which results in

dysfunctional ER retention of KDEL/RDEL-proteins, including Hsp47. To direct Hsp47 towards secretion in absence of BFA treatment, we deleted the "RVEL" sequence of Hsp47 in one of the constructs. Consistent with this concept, transfection of this construct results in increased secretion of Hsp47 protein and a higher (total) level of protection of EC from CIK, than mediated by the mostly ER retained form which only becomes transported to the cell surface via co-transport with collagen types I and IV (Hughes *et al.*, *Eur. J. Biochem.* 163:57-65 (1987)). The expression of Hsp47 and procollagen I were found to be tightly linked (Clarke *et al.*, *J. Cell Biol.* 121:193-199 (1993)).

Further PCR directed cloning of hsp47 into both pro- and eukaryotic protein expression vectors using eGFP-protein fusion tags enabled us to perform FACS analysis with full-length and truncational mutant *hsp47* gene cassettes. FACS analysis with full-length eGFP-Hsp47 protein stained a 26% subset of mature CIK.

Comparative analysis of the deduced peptide sequence of the human hsp47 gene and the available published data cited above (Clarke, *et al.*, *Biochem Biophys Acta* 1129:246-248 (1992)), indicate the presence of at least three distinct functional protein domains. We found one additional domain presented in this work. Hsp47 can be thus be depicted in a simplified fashion as a four domain protein (see Figure 15A). Through PCR with staggered primers and subsequent cloning into pro-and eukaryotic protein expression vectors, the protective function of truncated forms of Hsp47 was compared. See Figure 15C. The truncations were chosen to delete *hsp47* -gene fragments according to its presumed domain structure. See Figure 15B.

By sequence homology, Hsp47 is a serine protease inhibitor with theoretical specificity for lysine at the active site of incoming serine proteases. Serpins act as "bait" representing cleavable substrates which form a stable instead of a transitory covalent bond with the active center of serine proteases. Those then do not become released any more, thus blocking the active site of the serine protease they reacted with. Thus, serpins inactivate serine proteases in a stoichiometric ratio of 1:1. Despite the sequence relationship, no substrate

serine protease has so far been identified for Hsp47. Some authors expect the serpin-domain to be nonfunctional because its "active site" amino acids are modified (Hirayoshi *et al.*, *Mol Cell Biol* 11:4036-4044 (1991)). We mutagenized Hsp47 in truncational analysis across the serpin domain to evaluate whether a highly specific serine protease of the granule contents of CIK like granzyme A, but were not able to demonstrate a function of Hsp47 as an irreversible inhibitor of attacking granzymes in BLT esterase or SDS-PAGE gelshift assays (data not shown). Our data is paralleled by work with purified murine Hsp47 which failed to inhibit major serine proteases in BLT esterase assays in vitro (Davids *et al.*, *Bioorganic Chemistry* 23:437-438 (1995)). In <sup>51</sup>Cr-release assays, deletion of the serpin domain led to loss of function of the affected truncation mutants. However, as further deletions of the gene re-established full function to the smaller truncated Hsp47 mutants, expressing then little more than the H1,A-A2-consensus region discussed below, we assume that the loss of function of the C-terminal truncation of Hsp47's serpin domain causes a conformational change in the protein which normalizes with further truncation.

The second domain is the short C-terminal ER retention signal "RDEL." which we discussed above.

The third domain is the functionally evident collagen/RGD binding domain. So far it has not been directly localized. Truncation of either the C-terminal 32 amino acids or the N-terminal 34 amino acids of murine Hsp47 reportedly reduce the gelatin binding activity of mHsp47 (Davids *et al.*, *Bioorganic Chemistry* 23:437-438 (1995)). It is however not clear at present, whether these effects are secondary to conformational changes in the Hsp47-truncational mutants or whether they indicate the exact location of the functional RGD-binding domain. Significant sequence homology of mHsp47 exists with the crystallized collagen binding human protein C inhibitor (hPCI). This allowed for one attempt to theoretically model the three dimensional structure of Hsp47 and to predict a RGD binding loop at the C-terminal end of the protein (Davids *et al.*, *Bioorganic Chemistry* 23:437-438 (1995))

The fourth domain contains an alpha helical stretch closely resembling the peptide binding groove flanking  $\alpha_2$ -helix of the  $\alpha_2$ -domain of HLA-A2 and related MHC I proteins. This HLA-A2 consensus peptide was synthesized. This peptide is capable of protecting EC from CIK mediated killing in a dose responsive fashion (Figure 12). An immunoprotective domain of Hsp47 therefore contains a relatively small peptide domain which by itself is capable and sufficient to account for the protection of EC from non-MHC restricted killing.

We postulate that Hsp47 and the deduced peptide mediate protection of EC through a non-kill signal according to the paradigm of MHC I and the "missing self hypothesis" (Karre, *Semin Cancer Biol* 2:295-309 (1991); Karre *et al.*, *Nature* 319:675-678 (1986); Ljunggren *et al.*, *Immunol Today* 11:237-244 (1990)). Such a relationship has been described for killer inhibitory receptor (KIR) pathways (Höglund *et al.*, *Immunological Reviews* 155:11-28 (1997)). It appears possible that a non-kill signal is elicited in CIK cells through a yet to be identified counterreceptor recognizing qualitative and/or quantitative presence of the collagen chaperone Hsp47 in the vicinity of collagen deposition. Such a model could allow for a mechanistic explanation of tumor surveillance by CIK and other activated NK and T-cell populations *in-vivo*: Hsp47 is co-secreted as collagen chaperone by all cells actively depositing extracellular matrix containing collagens I or IV (Brewer *et al.*, *Embo J.* 16:7207-7216 (1997)). This secreted Hsp47 can bind to ubiquitous fibronectin consensus sequences (RGD) on fibronectin, osteonectin heparin and collagen molecules in the vicinity of the cell surface of the secreting cell (Nakai *et al.*, *Biochem Biophys Res Comm* 164:259-264 (1989)). Very recently it was reported that Hsp47 binds to tetraspanin CD9 on the surface of cells (Hebert *et al.*, *J. Cell Biochem* 73:248-258 (1999)). Tetraspanins are important membrane signal transducing molecules involved in the shaping of the T-cell repertoire (Levy *et al.*, *Annu Rev Immunol* 16:89-109 (1998)). A related tetraspanin, CD82 was recently reported to bind MHC class I molecules and to interfere with the KIR mediated non-kill signals of NK cells (Lagaudriere-Gesbert *et al.*,

*J. Immunol* 158:2790-2797 (1997)). Lack of adequate MHC I expression level leads to killing by default according to the "missing self hypothesis" in the context of NK target surveillance. In a similar fashion, lack of adequate collagen production by metastatic cells and their vasculature lacking contributory extracellular matrix (EM) production to their tissue of metastasis could thus lead to a lack of non-kill signal mediated by Hsp47 in the context of CIK, NK and T-cell mediated metastatic tumor surveillance.

Similarly, the physiologically aberrant vascular beds of tumors are characterized by fenestrated basal membranes and lack of normal EM production. In healthy individuals, only the ellipsoid meshwork of the spleen and liver have EC with fenestrated basal membranes. Those EC anchoring on physiologically fenestrated basal membranes are reported to have unusually high Hsp47 amounts (Kasai *et al.*, *Cell Tissue Res* 281:135-141 (1995)). The higher CIK cytotoxicity towards freshly confluent versus 5-day confluent EC *in-vitro* may represent lack of surface localized Hsp47 non-kill signal secondary to still incompletely laid down basement membrane after freshly seeding of TC flasks (Sauk *et al.*, *Biochem Biophys Res Comm* 172:135-142 (1990)). A Hsp47 mediated non-kill signal theory could provide a mechanism for CIK mediated *in-vivo* screening of tissue resident cells for extracellular matrix production and collagen production. According to this theory, detection and destruction of metastatic solid tumor cells would be the result from the latter lacking collagen and Hsp47 production. Additional support to this concept comes from reports that Hsp47 is down-regulated in cells undergoing oncogenic transformation (Clarke *et al.*, *J. Cell Biol* 121:193-199 (1993); Hirayoshi *et al.*, *Mol Cell Biol* 11:4036-4044 (1991)). In *in-vivo* studies, comparing a wide range of solid tumors, it was further found that Hsp47 becomes increasingly less expressed, as those tumors become more malignant and screening for Hsp47 levels was suggested to clinically follow tumor progression in cancer patients (Morino *et al.*, *In Vivo* 8:285-288 (1994)). Metastatic tumor nodules and their Hsp47 deprived vicinity thus could represent prime targets for cellular *in-vivo* tumor surveillance. EC lining the

fenestrated basal membrane found only in tumor neo-angiogenic vasculature represent by itself an additional target leading to specific attack and subsequent necrosis of tumor vascularization. Our SCID/hu-melanoma CIK results are fully consistent with this theory.

5           Blocking antibodies to KiR or p40 do not enhance the killing of tumor or endothelial targets. The involvement of other killer inhibitory receptors however remains possible.

          Previous studies demonstrated by FACS analysis that >95% of bulk generated CIK express CD3,  $TcR\alpha/\beta > TcR\gamma/\delta$  and  $CD8 > CD4$  (Lu *et al.*, *J. Immunol* 153:1687-1696 (1994)). Blocking antibody studies demonstrate that  
10           none of these molecules are functionally involved in the cytotoxicity against either EC or tumor targets (Table 2).

TABLE 2

CIK Cytotoxicity Blocking Studies with Monoclonal Antibodies

	Molecule Class:	Antibody Against:	> 15% Reduction of EC Lysis	> 15% Reduction of Tumor Lysis
5	Control	Irrelevant Isotype	-	-
	MHC	MHC I (W6/32)	-	-
	T-Cell			
	Markers:	TCR $\alpha/\beta$	-	-
		TCR $\gamma/\delta$	-	-
10		CD 2 (LFA-2)	-	-
		CD 4	-	-
		CD 8	-	-
	NK-Cell			
	Markers:	CD 16	-	-
15		CD 56	-	-
		p40 (via MAb NKTA 255)	-	-
		NKB1-Kir (via MAb DX 9)	-	-
	Cell Adhesion Molecules,			
20	$\beta 2$ intergrins	ICAM-1 (CD 54)	-	+
		LFA-1 (CD 18/11a)	-	+
		CD18 (LFA-1 $\beta$ -chain)	-	+
		CD 11a (LFA-1 $\alpha$ -chain)	-	+
		CD 11b (Mac-1 $\alpha$ -chain)	-	-
25		CD 11c (CR 4 $\alpha$ -chain)	-	-
	Other CAMs	CD 34	-	-
		PECAM-1 (CD 31)	-	-
		VAP-1	-	-
		N-Cadherin	-	-

30           Blocking Mab NK-B1 and NKp40 were also tested in  $^{51}\text{Cr}$  release cytotoxicity assays. Functional involvement of either molecule in CIK mediated cytotoxicity against tumor or EC targets has not been shown. These

receptors were assayed because expression of these NK markers is reported to occur on non-classical MHC I restricted cytotoxic T-cell (D'Andrea *et al.*, *J. Immunol* 155:2306-2310 (1995); Gumperz *et al.*, *J. Exp. Med.* 181:1133-1144 (1995); Lanier *et al.*, *J. Immunol* 153:2417-2428 (1994); Lanier *et al.*, *Immunol Today* 17:86-91 (1996); Phillips *et al.*, *Immunity* 5:163-172 (1996); Raulet *et al.*, *Cell* 82:697-700 (1995); Soderstrom *et al.*, *J. Immunol* 159:1072-1075 (1997)).

CIK can be used *in vivo* without co-injection of IL-2 (Lu *et al.*, *J. Immunol* 153:1687-1696 (1994)). This is a significant difference to past and current treatment of patients with IL-2 activated CD56<sup>+</sup> cells, IL-2 alone (Yang, *Cancer* 76:687-694 (1995)) or a combination thereof (Phillips, *Journal of Clinical Oncology* 5:1933-194 (1987)). Whereas the latter approaches reported a significant risk for the development of potentially fatal vascular leak syndrome (VLS), VLS was not observed in the *in-vivo* tumor purging models with SCID/hu mice. Nor was there significant CIK infiltrates in grafted normal human skin.

The reliable expansion and generation of highly specific anti-tumor activity is a sine qua non for any adoptive immune therapy cell production intended for clinical use. High dose Th<sub>1</sub>-type hormonal stimulation of  $\alpha/\beta$  T-cells with IFN- $\gamma$ , OKT-3 and IL-2 results in CIK AI cells with superior cytotoxicity against malignant hematopoietic and solid tumor targets. The use of turbine agitated bioreactors proves to be a plus, resulting in rapid, safe and improved selective expansion of the cytolytic CD3<sup>+</sup>56<sup>+</sup> subset. Yields of 10<sup>11</sup> cells are routinely obtained from a single buffy coat source. This is a sufficient number of effector cells for high dose and/or repeat treatment schedules in their clinical application. CIK can be generated from pure T-cell sources, maintained in long-term culture, cryo-preserved, and applied *in-vitro* and *in-vivo* without co-administration of IL-2. These characteristics make CIK superior choices for adoptive immune therapy of cancer as compared to LAK NK derivatives.

### Example 1



In vitro Cell CultureGeneration and Maintenance of CIK cells:

Whole venous blood from healthy community donors, umbilical cord blood and buffy coats obtained by the Stanford Blood Center served as source of peripheral blood lymphocytes (PBL). PBL cells were isolated using Ficoll-Hypaque (Pharmacia Fine Chemicals, Uppsala, Sweden) density gradient centrifugation. PBL were resuspended in RPMI 1640 (GIBCO-BRL/Life Technologies, Grand Island, NY) containing 50  $\mu$ m  $\beta$ -mercaptoethanol (ME), 100 IU penicillin-G ml<sup>-1</sup>, 100 IU streptomycin ml<sup>-1</sup> and 10% FCS (all: Sigma Chemical Co., St.Louis, MO) at a density of 0.5-2x10<sup>6</sup> cells/ml. Enriched populations of large granular lymphocytes (LGL) and T-cells were obtained by subsequent exclusion of plastic and nylon wool adherent cells. Source LGL were cultured in a humidified incubator with 5% carbon dioxide at 37 °C. Hormonal stimulation consisted of addition of recombinant human interferon gamma (rhu g-IFN) (a kind gift of Genentech, South San Francisco, CA) at 1000 IU ml<sup>-1</sup> at the start of culture day d0, subsequent stimulation on day one (d1) with soluble anti-human CD3 monoclonal antibody OKT-3 (derived from hybridomas by the American Tissue Culture Collection, Rockville, MD) at 50 ng per ml and recombinant human interleukin-2 (rhuIL-2) (a kind gift of Cetus/Chiron, Emeryville, CA) at 300-500 IU per ml. Cultures were maintained by addition of 50% fresh medium and 300-500 IU rhuIL-2 every 3-4 days for 21 to 28 days. Lytic activity was verified in <sup>51</sup>Cr release cytotoxicity assays against tumor and EC targets at day 14. Preferential expansion of the CIK CD3<sup>+</sup>, CD56<sup>+</sup> double positive phenotype was seen as early as day 10 in culture where an average of 12%-15% of the cells were double positive. Early CIK cultures could be further enriched by via a CD56-magnetic bead column to generate cultures which averaged 60% CD3<sup>+</sup> CD56<sup>+</sup> double positive phenotype (see Figure 2). For single cell CIK cloning T cells were purified from source LGL by FACS sorting for presence of CD3 or CD56 and absence of CD16 as described below. Cells were sorted at a mean density of 0.6 cells per well in a

96 well plate and expanded by co-culture with twice irradiated autologous feeder CIK at  $0.5 \times 10^6$  cells per ml with identical hormonal stimulation as described for bulk culture. Cells were expanded to more than  $2 \times 10^6$  cells each and then tested for clonal cytotoxicity as described below.

5 For large scale production of single donor CIK, healthy donor apheresis products were treated as above to isolate LGL. LGL were cultured at  $37^\circ\text{C}$  with identical hormonal stimulation as above, but at a tenfold density of  $5\text{--}10 \times 10^6$  cells/ml in turbine/impeller agitated bioreactors of 500 ml, 1,000 ml (Nalgene, Rochester, NY) and pO<sub>2</sub> and pH controlled 15,000 ml (Chemap, 10 Völketsvil, Switzerland) capacity. Use of 15L bioreactors allowed expansion up to  $3 \times 10^{11}$  cells from a single donor with the expansion of the CD3<sup>+</sup> CD56<sup>+</sup> cells exceeding 600 fold.

#### EC Culture, Generation and Maintenance of EC:

Freshly isolated single donor human umbilical chord endothelial cells (HUVEC) were a kind gift from Drs. J. Murphy and J. Alvarnas, (Stanford 15 University Div. Hematology. Stanford, CA). Pooled donor HUVEC were commercially obtained (ATCC, Rockville, MD). HUVEC were cultured on gelatin (Sigma Chemical Co., St. Louis, MO) coated tissue culture flasks, 6 and 24 well plates (Corning, Corning, NY) in media TC-199 or Ham's F12-K, each 20 supplemented with 100 IU penicillin-G per ml, 100 IU streptomycin per ml, 5 µg porcine heparin per ml, 20% FCS (all: Sigma Chemical Co., St. Louis, MO), and 80 µg endothelial mitogen per ml (Biomedical Technology Inc., Stoughton, MA) in a humidified incubator with 5% carbon dioxide at  $37^\circ\text{C}$ . Upon reaching 90% confluence HUVEC were passaged using 0.25% trypsin 25 and 0.02% EDTA in Ca<sup>2+</sup>+Mg<sup>2+</sup> free phosphate buffered saline (PBS).

Culture of EC on microcarrier beads was performed on 90 µm and 150 µm diameter spherical collagen coated acrylic cytocarriers (Kontes, Vineland, NJ) in magnetic stir bar agitated media TC-199 or Hams-F12K, each supplemented with 100 IU penicillin-G per ml, 100 IU streptomycin per ml, 5 µg porcine 30 heparin per ml and 20% FCS (all: Sigma Chemical Co., St. Louis, MO), and 80

µg endothelial mitogen per ml (Biomedical Technology Inc., Stoughton, MA) in a humidified incubator with 5% carbon dioxide at 37 °C.

Generation and Maintenance Tumor Target Cell Lines:

5 The hematopoietic tumor cell lines SU-DHL, OCI-Ly8, K562 and lymphoblastoid cell lines AMK (a kind gift of Dr. A. Krensky, Stanford University, CA) and RDL (a kind gift of Dr. R. Lopez, Stanford University, CA) were cultured in RPMI 1640 medium (GIBCO-BRL/Life Technologies, Grand Island, NY) containing 50 µm β-mercapto-ethanol (ME), 100 IU penicillin-G per ml, 100 IU streptomycin per ml and 10% FCS (all: Sigma  
10 Chemical Co., St.Louis, MO) at a density of 0.5-2x10<sup>6</sup> cells per ml in a humidified incubator with 5% carbon dioxide at 37 °C.

Melanoma cell lines WM 9 and 1205 LU (a kind gift of Dr. M. Herlyn, Wistar Institute of Anatomy and Biology, Philadelphia, PA) were maintained on tissue culture treated flasks (Beckton Dickinson, San Jose, CA) in MCDB  
15 153 medium (Life Technologies, Grand Island, NY) supplemented with 5 µg per ml insulin and 2% FCS in a humidified incubator with 5% carbon dioxide at 37 °C.

The constitutently cytoplasmic luciferase expressing cervical carcinoma cell line HeLaluc (a kind gift of C. Contag, Stanford University, Stanford, CA)  
20 was cultured in DMEM medium (GIBCO-BRL/Life Technologies, Grand Island, NY) containing 50 µm β-mercapto-ethanol (ME), 100 IU penicillin-G per ml, 100 IU streptomycin per ml and 10% FCS (all: Sigma Chemical Co., St.Louis, MO) in a humidified incubator with 5% carbon dioxide at 37 °C.

CTL cell-line AJY and specific CTL target cell-line AJ (lymphoblastoid)  
25 were kindly provided by the laboratory of A. Krensky (Stanford University, Stanford, CA).

**Example 2**Cloning and Expression of the Hsp47 Gene

Partial huHsp47 gene fragments derived by RT-PCR (a kind gift of Drs. Sanwal, Ontario, Canada) were amplified *in vitro* and artificial cloning sites  
5 introduced by PCR. These fragments correspond to nucleotides in Figure 1A. The nucleic acid encoding the amino terminal 39 amino acids of Hsp47 (excluding the signal sequence and first amino acid of the mature protein) was amplified with the following primers: 5' primer  
ACGTTTGGATCCAGGTGAAGA, 3' primer GTCCTTGGCCAT. The 5'  
10 primer incorporated a Bam HI site to facilitate cloning into further vectors. The 3' primer incorporated an Mlu NI site to facilitate fusing the nucleic acids encoding the amino and the carboxy terminal portion of the protein. The nucleic acid encoding the carboxy terminal 360 amino acids was amplified with the following primers: 5' primer  
15 GCAATGGCCAAGGACCAGGCAGTGGAG, 3' primer  
ACGCTCTGCTCAATATCCTTAAGTCTA. The 5' primer incorporated an Mlu NI site to facilitate fusing the two portions of the gene into one continuous reading frame and the 3' primer incorporated an Eco RI site to facilitate cloning into further vectors. Standard PCR conditions generally known to those of skill  
20 in the art were used to amplify the two fragments from the starting clones (see Figure 8). The resulting amplification products were digested with Mlu NI, purified and ligated to each other. The resulting nucleic acid was further digested with Bam HI and Eco RI and the huHsp47 gene cassette was directionally cloned using standard recombinant DNA techniques into pUC 19  
25 (Pharmacia, Uppsala, Sweden) which was used for construction purposes. The resulting plasmid, pUC/huHsp47, was amplified in recA bacterial host strain DH5a (Life Technologies, NY). The sequence of the cloned gene cassette was verified via fluorescent sequencing on an ABI sequencer (ABI,) using T7 DNA polymerase (Pharmacia, Uppsala, Sweden).

The huHsp47 gene cassette was PCR amplified using an alternative set of primers which results in a BamHI-EcoRI expressible gene cassette for bacterial protein expression. The expressible gene cassette was cloned into the pGEX 4T<sub>1</sub> vector (Pharmacia, Uppsala, Sweden), which produces a GST-fusion protein under the control of the IPTG inducible laq<sup>q</sup> promoter, and grown in the bacterial host strain BL21(DE3) (Stratagene, La Jolla, CA).

Expression of recombinant GST-fusion protein in bacterial host cells was induced with 20 mM IPTG via the lactose promoter laq<sup>q</sup> and overexpressed proteins purified via affinity chromatography on glutathione sepharose 4B (Pharmacia Biotech, Uppsala, Sweden), eluted at room temperature with 5-20 mM reduced glutathione in 50 mM Tris-HCl buffer at pH 8.0, analyzed quantitatively by Bradford assay and qualitatively by SDS-PAGE. The elution buffer was exchanged with PBS at pH 7.4 via PD-10 (Pharmacia Biotech, Uppsala, Sweden) gel filtration prior to use. When removal of the GST-fusion partner was desired, cleavage with 10 IU of thrombin (Sigma, St. Louis, MO) per µg fused GST moiety was performed for 10-12 hours at 37 °C using a thrombin-cleavage site present in the linker amino acids between GST and the recombinant protein of interest. Routinely 5-10 mg of recombinant huHsp47 fusion protein were obtained per 4 liters of BL21(DE3) host cells transformed by pGEX-4T-Hsp47 via glutathione affinity FPLC. Purified huHsp47 was used in <sup>51</sup>CR release cytotoxicity assays described above.

To dissect the role each domain of huHsp47 plays in protecting EC from CIK induced cytotoxicity, domain specific deletion mutants of huHsp47 were generated via PCR with nested primers based on the sequence disclosed in Figure 1 and previously published huHsp47 sequences. HuHsp 47 has at least four distinct functional domains. Beginning at the amino-terminus of the protein they are: a collagen/RGD binding domain, a domain with homology to the α-2 domain of human HLA-A2 molecules, a serine protease inhibitor (serpin) domain and an ER retention signal RDEL domain. Three specific deletions were generated. Deletion 1 removed the RDEL domain, deletion 2 removed the RDEL and serpin domains and deletion 3 removed the carboxy-

terminal 150 amino acids including the serpin and RDEL domains. The same 5' primer was used in PCR reactions to generate all three deletion mutants. The 5' primer sequence is: CGGAATTCTGGCCGAGGTGAAGAAACC. The 3' primer used to generate the RDEL deletion mutant was the same primer used to generate the huHsp47-GFP fusion protein and has the sequence:

AGTTCCCACTGTTCTACGACCTAGGGC. The deletion 2 3' primer is:

AACTCAACCTGTGTCTAGACCTATGGGC. The deletion 3 3' primer is:

ACGCGCTGCTCCTCCACGACCTAGGGC (see Figure 14). The 5' and 3'

primers incorporated BamHI and EcoRI restriction enzyme sites respectively to

facilitate subsequent cloning steps. The 5' and individual 3' primers were

mixed with the pUC/huHsp47 plasmid containing the huHsp47 gene cassette

and nucleic acids encoding each deletion mutant were PCR amplified in

separate reactions. The PCR products were purified, digested with BamHI and

EcoRI and ligated into either the pGEX-4T<sub>1</sub> vector to create huHsp47deletion-

GST fusion proteins for cytotoxicity assays (see below) or pEGFP-N1, which

also contained the mel secretion signal described above, to create eGFP-

Hsp47deletion fusion proteins for use as FACS probes.

Northern analysis of huHsp47 RNA expression was carried out via

$\alpha^{32}\text{P}$ -ATP nick translation labeling of the agarose gel purified full-length

huHSP47 cDNA as probe and human beta action (a kind gift of Dr. S.N.

Cohen) cDNA as internal control probe. Total RNA was harvested at 0 hour, 2

hour and 6 hour time points from EC undergoing BFA treatment. RNA for

each time point was purified, denatured, subjected to formamide flatbed

agarose gel electrophoresis, transferred to nitrocellulose, prehybridized,

hybridized with the  $^{32}\text{P}$ -labelled probes, washed under stringent conditions and

autoradiographed without intensifier screens at -80 °C using techniques well

known to those skilled in the art (see Sambrook et al. Molecular Cloning, A

Laboratory Manual, second edition, Cold Spring Harbor Laboratory Press,

1989). Results indicate that BFA treatment does not affect the expression of

the housekeeping gene beta action, but that BFA treatment does upregulate

huHsp47 gene expression over the period evaluated. See Figure 10. These

data are consistent with the observed increase in Hsp47 protein seen in BFA treated cells and isolated by immunoprecipitation experiments. See Figure 5.

Protease inhibitor free production of rhuHsp47 for granzyme A assays was carried out by growing protease free transformed BL21(DE3) bacteria in amp selective NZCYM medium using casein acidic hydrolysate in lieu of tryptic casein amino acids as nutritional peptide source for the host bacteria undergoing IPTG induction. To minimize incubation times, protein extracts were batch incubated on glutathione resin and initial removal of non GST proteins was carried out with a modified spin column approach. The remainder of the protocol was unchanged, thrombin was not utilized.

### Example 3

#### Cytotoxicity Assays

##### <sup>51</sup>Cr-Release Assay

Target cells were metabolically labelled with <sup>51</sup>Cr (Dupont- New England Nuclear, Boston, MA) by incubating 1x10<sup>6</sup> cells in 300 mCi <sup>51</sup>Cr at 37 ° C for 1-1.5 hours. The labelled cells were washed three times with phosphate buffered saline (PBS) containing 0.1% bovine serum albumin. The labelled cells were distributed in flat-bottomed 96 well microtiter plates at a concentration of 2x10<sup>4</sup> cells/well in triplicate. Effector (CIK) cells were added at the indicated ratios. Mabs were added prior to the addition of effector cells, and incubated for 15-30 minutes at room temperature. The final volume of the assay mixture in each well was 0.2 ml. After 4 hours at 37° C, the cells were collected by centrifugation and an aliquot of the supernatant was counted in a gamma counter (Micromedic Systems, Horsham, PA). The percentage of specific <sup>51</sup>Cr release was calculated according to the following equation:

$$\% \text{ specific } ^{51}\text{Cr release} = \frac{(\text{test release}) - (\text{spontaneous release})}{(\text{maximal release}) - (\text{spontaneous release})} \times 100\%$$

Spontaneous release was obtained by incubating the labelled cells in complete medium alone and maximal release by treatment of the cells with 1% NP-40.

The anti-tumor activity of adult and umbilical cord derived CIK cells were compared in cytotoxicity assays. The cytotoxicity results of the CIK cells derived from both sources were superimposable against B-cell lymphomas OCI-Ly8 and SU-DHL4/LAM53 and myeloid leukemia target K562. Both sets of CIK cells displayed marginal cytotoxicity against melanoma WM9 *in vitro*. Both sets of CIK cells were cytotoxic against freshly confluent HUVEC and less so against EC monolayers confluent for longer periods of time (see Figure 3).

Figures 3A, 3B and 3C describe 4 hour  $^{51}\text{Cr}$  release cytotoxicity assays with  $d_{21}$  CIK and various targets. CIK were derived from PBMC by culture in complete RPMI supplemented by the following: At  $d_0$ :  $\text{IFN}_\gamma$  at  $1000 \text{ u ml}^{-1}$ . At  $d_1$ : 5 ng soluble OKT-3 and  $500 \text{ u IL-2 ml}^{-1}$ . At  $d_5, d_9, d_{13}, d_{17}$  and  $d_{21}$ :  $500 \text{ u IL-2 ml}^{-1}$ . All cells were in a humidified incubator with 5%  $\text{CO}_2$  at  $37^\circ\text{C}$ . The hematopoietic tumor lines K562, OCI-Ly8, and SU-DHL4 were maintained in complete RPMI medium; melanoma line W9 was cultured in MCDB 153 medium supplemented with  $5 \text{ } \mu\text{g ml}^{-1}$  insulin and 2% FCS. HUV-EC were obtained from the ATCC. They were grown on 2% gelatin coated 24 well plates in TC-199 or Ham's F12-K medium, each supplemented with 20% FCS,  $5 \text{ } \mu\text{g porcine heparin ml}^{-1}$  and  $80 \text{ } \mu\text{g endothelial mitogen ml}^{-1}$ . Routine tryptic passaging was carried out when HUVEC reached more than 80% confluency;  $^{51}\text{Cr}$  release assays were routinely performed at the time point of fresh confluency.

Figure 3A is a cytotoxicity profile of adult and cord blood CIK  $d_{21}$  CIK shows high cytotoxicity against B-cell lymphomas OCI-Ly8 and SU-DHL4 (blue), as well as erythromyeloid leukemia K562 (yellow). Moderate cytotoxicity exists against nodular melanoma line WM9 (red) and freshly confluent HUV-EC (dark green).

Figure 3B shows the effect of EC culture duration after reaching confluency on the EC sensitivity to  $d_{21}$  CIK lysis: freshly confluent EC (dark



green) are lysed significantly more readily than are EC cultured for 5 additional days after reaching confluency (light green).

In Figure 3C clones of CIK were generated via plating of a mean of "0.6 cells per well" of a 96 well plate; then expanded by co-culture with twice irradiated, autologous feeder cells at  $0.5 \times 10^6$  cells  $\text{ml}^{-1}$ , and with identical hormonal stimulation as described above for bulk cultures. After individual clones expanded to more than  $2 \times 10^6$  cells each, they were tested for clonal cytotoxicity profile at a E:T ratio of 5:1. Target specificity of CIK is determined at the clonal level.

Expanded CIK single cell clones were tested to determine if sub-populations of cells exhibited target specificity in cytotoxicity assays performed as described above. Target specificity was observed, was line specific and was directed against tumor cells only, EC cells only or against both targets, the latter being the most common. The 2D10 CIK cell line tested displayed cytotoxicity against the OCI-Ly8 tumor cell line but not the HUVEC cell line. The 5F7 CIK cell line displayed cytotoxicity against HUVEC cells but not OCI-Ly8 cells. The 2C10 CIK cell line was cytotoxic against both the OCI-Ly8 and HUVEC cells.

#### BFA Treatment of Targets:

For brefeldin A (BFA) treatment, freshly confluent HUVEC monolayers, nonadherent target cells or CIK cells at  $0.5\text{--}2 \times 10^6$  cells per ml were incubated with medium supplemented with 0.5-10  $\mu\text{g}$  of brefeldin A (Böhringer Mannheim GmbH, Mannheim, Germany) per ml medium and incubated for 2-6 hours in a humidified incubator with 5% carbon dioxide at 37 °C. BFA was dissolved at 100  $\mu\text{g}$  per ml in 70% ethanol and used fresh or stored up to 4 weeks at -20 °C. Prior to cytotoxicity assay the cells were rinsed three times with PBS, and fresh medium not containing BFA was added for the remainder of the assay (Figure 4) .

BFA treatment interferes with the intracellular vesicle transport from the endoplasmic reticulum to the proximal and intermediate golgi compartments

preventing glycosylation of proteins directed for secretion and active vesicle transport of mature proteins to the cell surface (Lippincott-Schwartz et al., 1990; Lippincott-Schwartz et al., 1991; Lippincott-Schwartz et al., 1989; Orci et al., 1991; Vogel et al., 1993). Brefeldin treatment of endothelial cells leads to up regulation and secretion of a 46.5 KDa protein inhibitor presently identified as Hsp47 by immunodetection with MAB SPA470. Figure 5 shows immunoprotection with Mab N6 of  $^{35}\text{S}$  and biotin labeled HUVEC extracts after 6 hours of BFA induction. Lanes: M = marker, 1 = wt, 2 = 10  $\mu\text{M}$ , 3 =  $\mu\text{M}$  BFA.

Figure 5A is a silver stained SDS-PAGE, 10% polyacrylamide gel. Specific detection of p46.5 occurs only at a protective BFA dose. Note heavy and light chains of N6 Mab IgG.

Figure 5B is an auto-radiograph of a 12% polyacrylamide gel SDS-PAGE transferred to an NC membrane. A 46.5 KDa  $^{35}\text{S}$  radio-labeled band is present only at 10  $\mu\text{M}$  BFA.

Figure 5C demonstrates the positive identification of p46.5: It is huHsp47, immuno-detection with Hsp47 specific Mab SPA 470. Hsp47 migrates in this 12% PAGE with identical  $M_r$  of 46,500 as the radio-labeled peptide. The protein bands of p46.5 and huHsp47 are superimposable. Note, that due to the use of mouse IgG Mab N6 for the IP and detection of the primary mouse anti-Hsp47 Mab SPA470 with secondary goat-anti mouse peroxidase Ab, the heavy and light chains of the N6 Mab are also detected in this ECL assay.

BFA treatment of targets was able to suppress the CIK mediated lysis of EC in a dose dependent manner but did not affect the ability of CIK cells to lyse tumor targets (Figure 4). Both HUVEC and OCI-Ly8 cells were incubated with BFA at concentrations of 0, 2.5, 5, 7.5 and 10  $\mu\text{M}$  for 6 hours as described above, washed, labeled with  $^{51}\text{Cr}$  and used in a cytotoxicity assay with  $d_{21}$  CIK cells at an E:T ratio of 20:1. Preincubation of EC targets with 2.5  $\mu\text{M}$  BFA reduced CIK mediated cytotoxicity by 40%. Larger doses reduced cytotoxicity

further and preincubation in 10  $\mu$ M BFA was sufficient to completely block CIK mediated lysis of HUVECs.

<sup>51</sup>Cr Release Cytotoxicity Inhibition Assays with Blocking Mabs:

5 Prior to addition of effector cells 5  $\mu$ g of purified, NaN<sub>3</sub> free monoclonal antibody (Mab) was added per ml of medium to the target cells and incubated at room temperature (RT) for 15 minutes. Effector cells were added directly to the treated targets. The remainder of the assay was performed as detailed above, at 37 °C in a humidified atmosphere of 5% CO<sub>2</sub>.

10 Mabs to MHC class I (W6/32), T-cell markers CD2 (LFA-2), CD4, CD8, TcR $\alpha/\beta$  OR  $\gamma/\delta$ , and NK markers CD16, CD56, NK-B1(NKTA255) and Nkp40 (DX 9) do not block the cytotoxicity of CIK cells against either tumor or EC targets. Mabs to ICAM-1 (CD54), and LFA-1 (CD 18/11<sub>a</sub>, LFA-1  $\beta$ -chain, LFA-1  $\alpha$ -chain) do block cytotoxicity of CIK cells against tumor cells but not EC targets. No other anti-CAM Mab tested (CR 4  $\alpha$ -chain, CD 34, CD 15 31, VAP-1 or N-Cadherin) blocked the cytotoxicity of CIK cells against tumor targets. None of the antibodies tested blocked the cytotoxicity of CIK cells against EC targets indicating that CIK cells interact differently with tumor targets (see Table 2).

<sup>51</sup>Cr Release Cytotoxicity Assays with Protein and Peptide Addition:

20 Subsequent to <sup>51</sup>Cr labelling but prior to addition of effector cells 10-200  $\mu$ g FPLC purified Hsp47 (Example 7) protein or 10 nM to 10  $\mu$ M HPLC purified peptide AVLSAEQLR (Example 8), both in sterile PBS at pH 7.4 and diluted to the final concentration with RPMI, were added to targets per ml of medium and incubated for 30 minutes at 37 °C in a humidified atmosphere of 25 5% CO<sub>2</sub>. In assays using purified protein, targets were subjected to three subsequent washes with 10 volumes of Ca<sup>2+</sup>, Mg<sup>2+</sup> and 5% FCS containing PBS prior to providing fresh standard culture medium to the target cells. In assays using purified peptide, the added peptide was present for the entire assay. The

remainder of the assay was performed as detailed above, at 37 °C in a humidified atmosphere of 5% CO<sub>2</sub>.

Incubation of EC with 20µgm of gelatin-affinity purified (described below) huHsp47 per 300 µl was sufficient to completely protect the target from CIK mediated lysis (see Figure 7) in an identical fashion to the protection observed with 6h, 10 µM BFA treatment described above (see Figure 7).

The addition of as little as 1ηM of huHsp47-GST protein was able to reduce EC lysis by 50 %. Pre-incubation with 10 µM of huHsp47-GST protein completely protects EC from CIK mediated lysis (see Figure 9). The peptide AVLSAEQLR was also able to protect EC from CIK mediated lysis in a dose dependent manner in assays where the E:T ration was 20:1(see Figure 12). Peptide at a concentration of 1 µM was able to reduce EC lysis by 50 % and peptide at a concentration of 10 µM was able to suppress CIK mediated lysis almost completely. The peptide was capable of reducing specific target lysis by less than 10% when used at 10 µM in a CIK-OCI-Ly8 tumor cell <sup>51</sup>Cr cytotoxicity assay with an E:T ratio of 20:1.

The protective effects of each of the huHsp47 deletion mutants (Fig. 15B) were also analysed. See Fig. 15C. Deletion 1 mutant protein, lacking the RDEL domain, reduced CIK mediated lysis by 40% indicating that the RDEL domain is not involved in protecting the EC from cytotoxicity. Deletion 2 mutant protein, lacking both the serpin and RDEL domains, only reduced CIK mediated lysis by 20% suggesting either the serpin domain is involved in protecting EC from cytotoxicity or the resulting protein is folded aberantly resulting in a less active form of the protein. Deletion 3 mutant protein, lacking the carboxy-terminal 150 amino acids, was as effective as deletion 1 mutant protein in protecting EC from CIK mediated lysis suggesting that the reduced protection seen with the deletion 2 mutant protein is due to an altered conformation of the protein and is not due to the serpin domain participating in the protection. The deletion 3 mutant protein retains both the collagen/RGD binding domain and the α2-HLA-A2 homology domain which contains the AVLSAEQLR peptide sequence.

Protection of Endothelial Cells by  
Transfection With an Hsp47 Protein Expression Vector

Nucleic acids encoding huHsp47 and fragments were cloned into eucaryotic expression vectors. A nucleic acid encoding a fragment of huHsp47, in which the carboxy-terminal RDEL amino acid sequence is deleted, was PCR amplified from the pUC/huHsp47 plasmid using the following primers: 5' primer CGGAATTCTGGCCGAGGTGAAGAAACC, 3' primer AGTTCCCACTGTTCTACGACCTAGGGC. The amplified product was ligated to the melittin secretion signal and Kozak sequences derived from pMel-Bac (Invitrogen, San Diego, CA) and the resulting fragment was cloned into the multiple cloning site of pEGFP-N1 (Clontech, Palo Alto, CA) using general techniques well known to those of skill in the art. The resulting plasmid, eGFP-Hsp47, was transfected into EC.

Nucleic acids encoding full length huHsp47 and huHsp47 $\delta$ RDEL were cloned into the pBK-CMV (Stratagene, La Jolla, CA) and expressed in EC. The pBK vector was digested with Nhe I and EcoRI to remove the Lac promoter, most of the MCS and a portion of the Lac Z gene. The mellitin secretion signal peptide and Kozak sequence were PCR amplified from the pMel-Bac vector as described above and ligated into the pBK vector adjacent to the CMV IE promoter. Gene cassettes encoding either full length huHsp47 or huHsp47 $\delta$ RDEL were ligated into the pBK vector adjacent to the the mel signal sequence and the resulting vectors were used to transform EC. The CMV-neo-mel-huHsp47fulllength and CMV-neo-mel-huHsp47 $\delta$ RDEL vectors were purified from host cell cultures containing each vector and used to transfect EC using a low dose lipofectamine (GIBCO/BRL, NY) transfection protocol. The transfected EC were then analysed in CIK cytotoxicity assays.

EC were also co-transfected with a combination of vectors; eGFP-Hsp47, CMV-neo-mel-huHsp47fulllength and pCMV-CD20 (a vector expressing CD-20 under the control of the CMV promoter, a kind gift of Dr. S.vHeuvel, Massachusetts General Hospital, Boston, MA) using the low dose

lipofectamine transfection protocol. Cells were subjected to FACS analysis 24 hours after co-transfection.

EC targets were used in cytotoxicity assays 24 hours after lipofectamine mediated transfection of either CMV-neo-mel-huHsp47fulllength or CMV-neo-mel-huHsp47 $\delta$ RDEL into previously susceptible EC. Mature d<sub>21</sub> CIK cells were used at an E:T ratio of 20:1. EC over-expressing full length huHsp47 showed a 25% reduction in the CIK mediated lysis. EC over-expressing huHsp47 $\delta$ RDEL were entirely resistant to CIK mediated lysis (see Figure 11). Deletion of the RDEL peptide, which serves as an endoplasmic reticulum (ER) retention signal, from huHsp47, allows the protein to be freely secreted and protects the EC targets expressing the protein from CIK mediated lysis.

#### Example 4

##### Fluorescence Activated Cell Sorting and Imaging

For Fluorescence Activated Cell Sorting (FACS) 10<sup>6</sup> effector or target cells were washed twice with Ca<sup>2+</sup>, Mg<sup>2+</sup> and 5% FCS containing PBS buffer, suspended in 50  $\mu$ l of this buffer and placed at 0-4 °C in the dark. 1  $\mu$ g of FITC-, PE-, Per-CP or TriColor labelled monoclonal antibody (Beckton Dickinson, San Jose, CA) per staining reaction of 10<sup>6</sup> cells was mixed with the cells and incubated for 15 to 30 minutes at 0-4 °C in the dark. Stained cells were subjected to three subsequent washes with 10 volumes of Ca<sup>2+</sup>, Mg<sup>2+</sup> and 5% FCS containing PBS and taken up in 500  $\mu$ l of this buffer for analysis or sort by FACScan/FACStar (Beckton Dickinson, San Jose, CA) with excitation at 488 nm and the appropriate channel filters for detection of emission.

Cell staining for immunofluorescence microscopy was carried out in a similar manner and stained cells analyzed with a Axiophot fluorescence camera system (Zeiss, Hanau, Germany).

The huHsp47 $\delta$ RDEL-enhanced green fluorescent protein (eGFP) fusion protein, expressed under the control of the CMV IE promoter, was isolated as described above and used as a FACS probe to search for huHsp47 cell surface binding proteins. FACS analysis of mature d<sub>21</sub>CIK with eGFP-tagged

rhHsp47 as molecular probe was carried out by first staining with PE-Mab and specifically PE-CD56 if desired, then by suspending the washed CIK in 10 µg eGFP-huHsp47 per ml of PBS, incubating for 15 minutes at 0-4 °C in the dark, removal of unbound protein by centrifugation in a refrigerated eppendorf  
5 centrifuge at 1000 rpm at 0-4 °C. Stained cell pellets were either resuspended in 500-1000 µl of PBS immediately prior to the FACS analysis or suspended in PBS fixans containing 3% (w/v) formaldehyde prior to analysis. After staining, cells were sorted using the FITC channel. 26% of the cells analysed were positive for both eGFP-huHsp47 and PE-CD56 probes indicating that a distinct  
10 subset of the CIK cells had receptors for huHsp47.

EC cells co-transfected with eGFP-Hsp47, CMV-neo-mel-huHsp47fulllength and pCMV-CD20 using the low dose lipofectamine transfection protocol were analysed 24 hours after co-transfection. Cells were stained with PE-CD20 Mab and analysed for the presence of both the CD-20  
15 antigen and eGFP-Hsp47 using the FITC channel. Over 79% of the cells were positive for both probes indicating that there was an extremely high transfection efficiency.

### Example 5

#### Native Protein Purification and Analysis

##### 20 Protein Extraction from Target Cells

Prior to cell harvest, targets were cultured or treated BFA as described above, and dead cells removed from EC monolayers by washes with  $\text{Ca}^{2+}$ ,  $\text{Mg}^{2+}$  containing PBS, from nonadherent targets via density gradient centrifugation. Cell lysis was performed with ultrasonic cell disruption (Branson, New  
25 Brunswick, NJ) in PBS buffer containing 1% Triton-X 100 (Tx-100, Sigma, St. Louis, MO) and complete protease inhibitor cocktail (Boehringer Mannheim, Mannheim, Germany).

For metabolic  $^{35}\text{S}$  methionine/cysteine labelling cells were cultured for 22 hours in cysteine and methionine free RPMI 1640 or Hams-F12K (Irvine

Scientific, Santa Ana, CA) media with dialyzed FCS, otherwise as described above, prior to the addition of 100  $\mu$ Ci ml<sup>-1</sup> of Translabel (PIERCE/Amersham, Amersham, UK) <sup>35</sup>S methionine/cysteine at the start of BFA, control treatment and incubated for the indicated times. Pulse-chase labelled proteins were analyzed by SDS-PAGE, western analysis or autoradiography and silver staining.

Labelling of membrane proteins by biotinylation was carried out after applicable treatments with BFA and subsequent stringent removal of dead cell. Water soluble, lipid insoluble 23-atom linker arm modified biotin analogues NHS-SS-biotin and NHS-LC-biotin (PIERCE, Rockford, IL) were added to a final of 1 mg ml<sup>-1</sup> of PBS and cell monolayers incubated for 30 minutes at room temperature. The reaction was stopped by removal of the biotin labelling reaction and three washes with 10 volumes of 100 mM Tris-HCl, 150 mM NaCl, 5 mM EDTA buffer at pH 7.4 providing excess amine to react with residual NHS-groups.

Purification of NHS-SS-biotin biotinylated proteins was carried out by affinity chromatography on a resin of streptavidin immobilized to 6% crosslinked agarose (Sigma, St. Louis, MO) under UV260nm control (Pharmacia, Uppsala, Sweden). Bound proteins were released from the streptavidin matrix by cleaving the disulfide bond of the 23 atom-SS-biotin linker arm with 100 mM  $\beta$ -mercapto ethanol: This results in elution of the previously biotin-tagged protein leaving the biotin moiety bound to the streptavidin resin. Specifically eluted proteins were analyzed by SDS-PAGE, autoradiography and silver staining. Detection of NHS-LC-biotinylated proteins was carried out by SDS-PAGE, subsequent transfer to solid support membranes and reaction with streptavidin-alkaline phosphatase (AP). Bound streptavidin-alkaline phosphatase was detected via colourimetric BCIP/NBT blot assay. Subsequent autoradiography was carried out with metabolically <sup>35</sup>S methionine/cysteine labelled extracts.



**Example 6**Generation and Preparation of AntibodiesGeneration and preparation of polyclonal antibodies:

5 Generation of polyclonal rabbit antisera was performed after harvest of non-immune/preserum from three female New Zealand rabbits. One mg of total protein was used per immunization day and animal. Ultracentrifugation derived wild-type (wt) and TM treated lymphoblastoid target cell membrane preparations were used as antigens. Antigens were prepared as emulsion in complete Freud's adjuvant (CFA, Sigma, St. Louis, MO) and injected  
10 subcutaneously. This was followed by three booster immunizations at six week intervals as above, but substituting FCA for Freud's incomplete adjuvant (FIA). Titer development was monitored by antigen spot assay of rabbit ear vein blood samples obtained at time points of boosting. Serum was separated from the cell product of collected specimen by coagulation and subsequent centrifugation,  
15 aspirated and stored frozen at -20 °C until used.

Preparation of Monoclonal Antibodies:

Limited production of monoclonal N6 antibody in-vivo: Pan-anti-heat shock protein Mab N6 hybridoma (a kind gift of Dr. K Nagata, Kyoto, Japan) was injected intraperitoneally into six-week old female Balb/c mice. Ascitis  
20 fluid was harvested, shockfrozen in liquid N2 and stored at -20 °C until needed.

Antibody Purification:

FPLC purification of polyclonal and monoclonal antibodies from serum or ascitis was performed after 7x dilution in PBS and chromatography on Protein A sepharose (Pharmacia, Uppsala, Sweden) with UV260nm control. Bound Ig  
25 fractions were eluted with 25 mM glycine buffer at pH 2.5 and rapidly titrated back to pH 7.4 by addition of TRIZMA (Sigma, St. Louis, MO) base. Protein amounts were determined by Bradford assay at 595 nm in a DU-650 spectrophotometer (Beckman, Palo Alto, CA).

Immunoprecipitation was carried out by first preparing preformed immune complexes as follows: Purified specific mouse monoclonal or rabbit polyclonal antibody were incubated in stoichiometric ratio with purified secondary goat anti-mouse or anti-rabbit antibody at a concentration of 10 µg ml<sup>-1</sup> of 150 mM NaCl, 10 mM Tris-HCl, 1% (w/v) NP40, pH 7.8 with gentle agitation for 3 hours at 4 °C. In a separate reaction their respective isotype matched control antibodies were coupled to protein A sepharose beads (Pharmacia, Uppsala, Sweden). The formed immune complexes were harvested by centrifugation at 13,000 rpm in a refrigerated Eppendorf centrifuge (Eppendorf, Hamburg, Germany) for 15 minutes at 4 °C, washed twice with one volume of 500 mM NaCl, 50 mM Tris-HCl, 2% BSA, 0.25% (w/v) SDS, 0.5% (w/v) Tx-100, pH 7.5 and taken up in 120 µl of 150 mM NaCl, 10 mM Tris-HCl, 1% (w/v) NP40 at pH 7.8. For protein extraction, 2x10<sup>7</sup> cells per ml of buffer were lysed by sonication in 150 mM NaCl, 10 mM Tris-HCl, 1% (w/v) NP40, 1 mM PMSF, 20 µg ml<sup>-1</sup> trypsin-inhibitor, 5 mM EDTA at pH 7.8. Cell debris was removed by centrifugation at 13,000 rpm for 30 minutes at 4 °C. The lysate was cleared with the isotype matched control antibodies coupled to protein A sepharose beads by agitation for 3 hours at 4 °C and subsequent removal of the beads by centrifugation at 13,000 rpm for 15 minutes at 4 °C. Immunoprecipitation with specific antibody was carried out by adding 30 µl of cleared protein extract to 200 µl of 150 mM NaCl, 10 mM Tris-HCl, 1% (w/v) NP40, pH 7.8 and incubating with 10 µl of preformed specific immune complexes per reaction with gentle agitation for 3 hours at 4 °C. The specific immune precipitate was recovered by centrifugation at 13,000 rpm for 30 minutes at 4 °C, washed twice with 10 mM Tris-HCl at pH 6.8 and analyzed by SDS-PAGE and subsequent silver stain, transfer to solid support and streptavidin-AP detection and autoradiography.

AMK targets were first treated for 16 hours with TM at 5 µg per ml or left untreated and then treated concurrently with BFA and <sup>35</sup>S met/cys (in the presence of TM in the originally treated group) as described above followed by a 2 hour chase. Crude cell extracts were immunoprecipitated (IP) either with

polyclonal antisera prepared against AMK membrane preparations described above or with pan-anti heat shock Mab N6. Autoradiographs of the SDS-PAGE separated IP with either antibody of total cellular protein shows the presence of a <sup>35</sup>S labeled 46.5 KDa protein in CIK resistant untreated lymphoblastoid cells which was absent in the TM treated cells.

EC targets were cultured in the presence of no BFA, the protective dose of 10 μM BFA, or the non-protective dose of 30 μM BFA concurrently with <sup>35</sup>S met/cys as described above. Samples were processed as described for the AMK cells. EC targets cultured in the presence of BFA also contained a <sup>35</sup>S labeled 46.5 KDa protein. This protein was not detectable in IP from EC cultured without BFA or with the non-protective dose of 30 μM BFA (see Figure 5).

This 46.5 KDa protein was identified as Hsp47 in a Western analysis using the Mab SPA-470 antibody. <sup>35</sup>S labeled HUVEC extracts prepared from BFA treated cells described above were immunoprecipitated as described using the N6 pan-heat shock Mab. Precipitates were run on a 12% SDS-PAGE, transferred to NC membrane, blocked and reacted with the SPA-470 antibody using techniques well known to those of skill in the art. The presence of the SPA-470 antibody bound to the 46.5 KDa protein and the presence of the N6 heavy and light chains were detected using goat-anti mouse horseradish peroxidase staining assay(ECL).

Biotinylation of live adherant EC with water soluble, lipid insoluble Biotin analog NHS-LC-biotin performed as described above followed by IP of cell extracts using either the anti-AMK polyclonal antibody or N6 Mab and subsequent SDS-PAGE, transfer to NC membrane and detection using a streptavidin-Alkaline phosphatase system showed that the Hsp47 protein induced by BFA treatment and a minor protein of 27 KDa are localized on the cell surface of EC targets.

**Example 7**Purification and Analysis of Hsp47 Proteins

Native and recombinant protein purification was performed by subjecting protein crude extracts to anion-exchange FPLC chromatography on Q-sepharose® at pH 7.4, and pH 6.8, eluting separated fractions by increasing ionic strength and subsequent size exclusion FPLC chromatography on Superose 12 separating fractions with increasing elution volumes  $V_e$ , controlled by a UV-M monitor 214, 260 and 280nm and recorded in analogue mode (all equipment and resins: Pharmacia, Uppsala, Sweden) (see Figure 6). Protein amounts were quantitated on-line by integration of the area under the recorded peak of absorbance (AuP) and by Bradford assay of sampled fractions. Qualitative analysis was performed by SDS-PAGE and subsequent Coomassie and silver staining, as well as membrane transfer and western blot, when specific antibodies were available.

Figure 6A is a  $UV_{260nm}$  profile of native Hsp47 purification from EC on Q-sepharose ff anion exchange resin at pH 8.9 175 flasks of confluent HUVEC were induced with 10  $\mu$ M BFA as described. In the first step of ion exchange purification, proteins were separated with the same resin at pH 7.4 and hsp47 containing fractions pooled. Subsequently the proteins were diluted five fold in low salt buffer at pH 8.9 to yield buffer A (75 mM Tris•Cl pH 8.9, 175 mM NaCl, 2 mM DTT, 0.5 mM EDTA, 2 mM PMSF), and loaded on the ion exchange column. Bound proteins were subjected to a linear salt gradient with buffer B = A+100 mM NaCl. Separation of individual protein peaks was enhanced by manually switched step salt concentration plateaus, based upon OD extinction.

Figure 6B is a 12% SDS-PAGE analysis with silver staining of the collected p46.5 containing fractions. Those were pooled and concentrated by YD10 membrane to 1.0 ml.

Figure 6C is a 12% SDS-PAGE analysis with silver staining of the collected p46.5 containing fraction after size exclusion FPLC chromatography on Superose 12. Yield: 240µg.

5 Affinity FPLC chromatography was performed by loading protein extracts of wt and BFA treated EC onto gelatin-sepharose (Pharmacia, Uppsala, Sweden) matrix with help of a peristaltic pump and incubating the extract for 30 minutes at 0-4 °C. Unbound and unspecifically bound proteins were removed by washing with 7 column volumes of high salt buffer followed by original protein extraction buffer until the UV260nm detection at AU 0.02  
10 returned to baseline. pH specific elution was carried out with PBS buffer at pH 5.5, alternative eluents included 100 mM ATP in PBS and 100 µM RGD peptide in PBS, both at pH 7.4.

Immunoaffinity FPLC chromatography was carried out in a similar manner, with the protein extract loaded onto a specific Mab column: 10 mg of  
15 purified Mab N6 was coupled via a covalent thioether bond with a 13 atom spacer arm to activated, crosslinked agarose via the Sulfolink (Pierce, Rockford, IL) protocol. Unspecifically bound proteins were removed from the columns as above, specifically immuno recognized proteins were eluted via pH 2.5 elution in 50 mM Na<sup>+</sup> citrate buffer.

20 Hsp47 may also be gelatin-affinity purification. 175 flasks of confluent HUVEC were induced with 10 µM BFA for 6 h at 37°C, 5% CO<sub>2</sub>, in F12K medium supplemented with 20% FCS and standard EC culture supplements. Subsequently, adherent EC were harvested by scraping in detergent buffer (75 mM Tris•Cl, 175 mM NaCl, 2 mM DTT, 0.5 mM EDTA, 0.5% Tx-100, 2 mM  
25 PMSF), lysed by sonification and the extract cleared from cell debris by centrifugation at 15,000 rpm in a refrigerated eppendorf centrifuge. The cleared extract was loaded onto a 5 ml covalently coupled gelatin 4B Sepharose column via peristaltic pump at 0.5 ml min<sup>-1</sup>. Unbound proteins were discarded. The column was then washed with detergent free buffer (as above, w/o Tx-  
30 100), low salt buffer (no NaCl) and high salt buffer (plus 500 mM NaCl) under UV<sub>260nm</sub> control until baseline readings were achieved. Specific elution of

Hsp47 was achieved by lowering the pH of the detergent free buffer to pH 4.8. Collected eluate was immediately back-titrated to pH 7.4. The column was immediately regenerated and washed with pH 7.4 buffer.

5 Figure 2 shows the results of incubation of EC with gelatin-affinity purified Hsp47 for 30 minutes at 21°C, prior to <sup>51</sup>Cr release CIK cytotoxicity assay. Excess non-bound Hsp47 protein was removed from the wells by three washes prior to the addition of CIK effector cells. As a control, untreated BFA induced HUVEC targets were compared to the protein treatment group. Human Hsp47, the protein of 46.5 KDa purified with pH specific elution from  
10 gelatin matrix, protects EC.

### Example 8

#### Hsp47 Peptides

##### Peptide Synthesis

15 A hydrophilic 9 mer peptide of the sequence AVLSAEQLR which corresponds to the region of shared homology of Hsp47 with HLA-A2 was synthesized by the Stanford PAN facility on a BioApplied® peptide synthesizer, HPLC purified and verified by amino acid sequencing. Pure peptide was lyophilized and stored at -20 °C. The peptide was soluble in aqueous solution and used in PBS and RPMI 1640 at indicated concentrations.

20 Amino acid sequencing of purified native p46 and proteolytic fragments derived from it was performed by A. Willis (Immuno Chemistry Unit of the MRC, Oxford University, Oxford, UK).

### Example 9

#### In-vivo Analysis

##### 25 SCID/hu Skin Graft-Melanoma-CIK Tumor Purging Model

Female 6 week old balbc/SCID mice were irradiated at 500 Rads at the day prior to skin grafting. Human full thickness foreskin grafts were prepared

by aseptically exposing their subcutaneous vascular beds, and trimming their edges to yield a 20x30 mm graft. Foreskins yielding only smaller grafts or older than 24 hours of shelf age were rejected. Anesthetized animals were prepared in a 37 °C heated operative environment by aseptically denuding matching areas on their flanks. The full-thickness skin grafts were placed with interrupted absorbable suture, aseptically bandaged and the grafted animals housed separately until full engraftment was evident. After engraftment,  $1 \times 10^5$  melanoma WN9 or 1205 LU cells in PBS were injected intradermally into a single site of the human full-thickness skin of triplicate animals and tumor growth monitored by observation and palpation. Animals with evident nodular tumor burden exceeding a diameter of 5 mm were injected intraperitoneally with  $1 \times 10^7$  CIK cells (suspended in RPMI without FCS or IL-2) of proven lytic activity in same day  $^{51}\text{Cr}$  release cytotoxicity assay at CIK culture day d14-21. Animals were sacrificed at 0, 24, 48 and 72 hours following CIK injection, internal organs and the human skin/melanoma graft fixed separately and analyzed by light microscopy of H&E and immuno-peroxidase anti-human CD3, human PE-CAM and human Kir antigens stained tissue sections. IP injected CIK cells selectively infiltrated the vascular beds of the melanoma tumor, which was confirmed by IPOX histologic immunodetection using anti-human PE-CAM Mab. Excluding a mild infiltrate in the portal area of the mouse liver, vasculature and skin tissue of the normal human skin graft or the murine host were not infiltrated by CIK *in vivo*.

SCID/hu HeLa Luciferase Hsp47 Secretion  
CIK Tumor Purging Live Imaging Model

Female 6 week old balbc/SCID were irradiated at 500 Rads at the day prior to intraperitoneal injection of  $2 \times 10^4$  HeLa<sub>luciferase</sub> cervical carcinoma cells capable of imagable in-vivo luminescence. The CIK treatment group were injected with CIK at a 20:1 E:T ratio ( $1 \times 10^6$  cells) via a tail vein injection at the time of the HeLa cell challenge. Those HeLa<sub>luciferase</sub> cells were transiently transformed 24 hours prior to injection via lipofectamine (GIBCO/BRL, NY)

protocol with a vector constitutively expressing full-length huHsp47 directed to secretion under immediate early CMV promoter control or respective irrelevant control vectors. Transfection efficiency was monitored via co-transfection of eGFP and the Hsp47 construct with subsequent fluorescence laser microscopy analysis of transformed HeLa<sub>luciferase</sub> monolayers with a Axiophot (Zeiss, Jena, Germany) fluorescence microscope described above. Injected animals were monitored at 0, 2 and 7 days for *in-vivo* chemo-luminescence of remaining HeLa<sub>luciferase</sub> post CIK challenge with a quantitative digital imaging process in collaboration with Dr. C. Contag (see Figure 13). Animals receiving the CIK injection demonstrated a dramatic decrease in live, light emitting tumor burden one week after CIK purging. Animals which did not receive the CIK cells demonstrated widespread IP growth of HeLa cells. The CIK cells demonstrated a high and rapid solid tumor purging potential.

#### Murine GvHDModel

Male Balb/c mice were twice irradiated with 500 Rads prior to transplantation. Bone marrow (BM) and spleen derived stem cells were obtained from B10D2 (H-2<sup>d</sup> background) mice.  $2.5 \times 10^6$  BM and  $1 \times 10^6$  spleen cells were pooled as a graft source for each recipient male Balb/c animal. This non-enriched stem cell population of cells were either suspended in RPMI containing 300  $\mu$ g Hsp47-Gst protein in 300  $\mu$ l total volume, or, in buffer containing 100  $\mu$ g of Gst control protein alone. Cells and protein were given in a single injection on day 0. No other immunosuppressive treatments were given. Animals were monitored for 4 weeks after the bone marrow transplant. All 5 animals receiving the Hsp-Gst protein treatment were protected from the onset of GvHD, fur ruffling, weight loss and death. Animals receiving only the Gst protein all exhibited signs of GvHD, fur ruffling and weight loss, and 4 of the 5 animals died (see Figure 15).



**Example 10****FACS Analysis**

This Example discloses the use of modified Hsp47 to identify and separate immune cell subsets.

5           A variety of immune cells respond to Hsp47 and do not attack endothelial cells. In order to identify such cells and to enrich adoptive immunotherapeutic populations from these non-endothelial reactive cells in the presence of Hsp47, fluorescent tagged Hsp47 can be used in a standard cell sorting apparatus to identify and separate such cells. Such use can also serve as a diagnostic tool  
10          for immune cell subset analysis in various disease states.

          eGFP-tagged Hsp47 FACS probe was generated through additional genetic engineering (Figure 8A). This enabled searching for both potential huHsp47 receptors and for CIK subsets expressing such Hsp47 binding proteins. FACS analysis of mature d<sub>21</sub> CIK with eGFP-tagged HuHsp47 as  
15          FACS probe for the FITC channel, with CE56-PE co-staining, is based on differences in the Hsp47 binding ability of the analyzed CIK. The eGFP tagged Hsp47 probe stained a quarter of mature CIK (26%).

          The foregoing description details specific methods which can be employed to practice the present invention. Having detailed such specific  
20          methods, those skilled in the art will well enough know how to devise alternative reliable methods at arriving at the same information in using the fruits of the present invention. Thus, however, detailed the foregoing may appear in text, it should not be construed as limiting the overall scope thereof; rather, the ambit of the present invention is to be determined only by the lawful  
25          construction of the appended claims. All documents cited herein are expressly incorporated by reference.

**WHAT IS CLAIMED IS:**

1. A method for reducing immune-mediated damage to cells, tissue or organs comprising contacting a tissue, organ or cell other than endothelial cells with an immunoprotective amount of an Hsp47-related immunoprotective polypeptide.  
5
2. The method of Claim 1 wherein said Hsp47-related immunoprotective polypeptide comprises the sequence AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AX<sub>4</sub>X<sub>5</sub>X<sub>6</sub>R wherein X<sub>1</sub> is V, L, A or T, X<sub>2</sub> is L or H, X<sub>3</sub> is S or V, X<sub>4</sub> is D or E, X<sub>5</sub> is Q, K or R, and X<sub>6</sub> is L or V.
3. The method of Claim 1 wherein said Hsp47-related immunoprotective polypeptide comprises the sequence AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AEQLR, where X<sub>1</sub>, X<sub>2</sub> and X<sub>3</sub> can be any amino acid.  
10
4. The method according to Claim 3 wherein X<sub>1</sub> is V or A, X<sub>2</sub> is L or H and X<sub>3</sub> is S or V.
5. The method according to Claim 1 wherein said polypeptide comprises the sequence AVLSAEQLR.  
15
6. The method according to Claim 1 wherein said immunemediated damage caused by an autoimmune disease, graft vs. host disease or host vs. graft disease.
7. The method according to Claim 1 wherein said contacting is *ex vivo*.
8. A method for reducing immune-mediated damage to cells or tissue comprising:  
20  
    contacting a tissue, organ or cells other than endothelial cells with an immunoprotecting amount of a composition comprising Hsp47 polypeptide.

9. The method according to Claim 8 wherein said Hsp47 polypeptide comprises an amino acid sequence selected from the group consisting of:
- (a) the amino acid sequence shown in Figure 1;
  - (b) an amino acid sequence having at least about 70% amino acid  
5 sequence identity with the amino acid sequence of (a);
  - (c) an immunoprotecting fragment of (a) or (b); and
  - (d) an immunoprotecting variant of (a) or (b).
10. The method according to Claim 9 wherein said immunoprotecting fragment of Hsp 47 comprises the amino acid sequence AVLSAEQLR.
- 10 11. The method according to Claim 9 wherein said immunoprotecting variant of Hsp 47 is a conserved variant of the sequence AVLSAEQLR.
12. The method according to Claim 9 wherein said immunoprotecting variant of Hsp 47 comprises the sequence AAHVAEQLR.
13. The method according to Claim 8 wherein said Hsp47 polypeptide  
15 comprises an amino acid sequence selected from the group consisting of:
- (a) the amino acid sequence shown in Figure 1;
  - (b) an amino acid sequence encoded by a nucleic acid that hybridizes  
to a nucleic acid that encodes the amino acid sequence shown in Figure 1;
  - (c) an immunoprotecting fragment of (a) or (b); and
  - 20 (d) an immunoprotecting variant of (a) or (b).
14. The method according to Claim 8 wherein said contacting is *in vivo*.
15. The method according to Claim 8 wherein said immune mediated damage is caused by an autoimmune disease, graft vs. host disease or host vs. graft disease.

16. A method for reducing immune-mediated damage to tissue, organ or cells comprising:  
contacting a tissue, organ or cells other than endothelial cells with an immunoprotecting amount of a composition comprising brefeldin.
- 5 17. The method according to Claim 16 wherein said immune mediated damage is caused by an autoimmune disease, graft *vs.* host disease or host *vs.* graft disease.
18. The method according to Claim 16 wherein said contacting is *ex vivo*.
19. A method for reducing immune-mediated damage to tissue, organs or  
10 cells comprising:  
contacting a tissue, organ or cell other than endothelial cells with an expressible nucleic acid encoding an Hsp47-related immunoprotective polypeptide.
20. The method according to Claim 19 wherein said Hsp47-related  
15 immunoprotective polypeptide comprises the sequence  $AX_1X_2X_3AX_4X_5X_6R$  wherein  $X_6$  is V, L, A or T,  $X_2$  is L or H,  $X_3$  is S or V,  $X_4$  is D or E,  $X_5$  is Q, K or R, and  $X_6$  is L or V.
21. The method according to Claim 19 wherein said 47-related immunoprotective polypeptide comprises the sequence  $AX_1X_2X_3AEQLR$ ,  
20 where  $X_1$ ,  $X_2$  and  $X_3$  can be any amino acid.
22. The method according to Claim 21 wherein  $X_1$  is V or A,  $X_2$  is L or H and  $X_3$  is S or V.
23. The method, according to Claim 19, wherein said polypeptide comprises the sequence AVLSAEQLR.

24. The method according to Claim 19, wherein said immune mediated damage caused by an autoimmune disease, graft vs. host disease or host vs. graft disease.
25. The method, according to Claim 19, wherein said contacting is *ex vivo*.
- 5 26. A method of adoptive immune therapy to treat cancer comprising:  
culturing T-lymphocytes from a donor under conditions wherein a specific subset of said T-lymphocytes are expanded; and  
introducing said expanded subset of said T-lymphocytes to a patient in conjunction with an immunoprotecting amount of an immunoprotective agent  
10 selected from the group consisting of brefeldin and Hsp47-related immunoprotective polypeptides.
27. The method according to Claim 26 wherein said immunoprotective polypeptide comprises the sequence AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AX<sub>4</sub>X<sub>5</sub>X<sub>6</sub>R wherein X<sub>1</sub> is V, L, A or T, X<sub>2</sub> is L or H, X<sub>3</sub> is S or V, X<sub>4</sub> is D or E, X<sub>5</sub> is Q, K or R, and X<sub>6</sub> is L or V.
- 15 28. The method of Claim 26 wherein said immunoprotective polypeptide comprises the sequence AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AEQLR, where X<sub>1</sub>, X<sub>2</sub> and X<sub>3</sub> can be any amino acid.
29. A method according to Claim 28 wherein X<sub>1</sub> is V or A, X<sub>2</sub> is L or H and X<sub>3</sub> is S or V.
- 20 30. A method for identifying cells which bind to a polypeptide comprising contacting said cells with a polypeptide comprising the sequence AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AX<sub>4</sub>X<sub>5</sub>X<sub>6</sub>R wherein X<sub>1</sub> is V, L, A or T, X<sub>2</sub> is L or H, X<sub>3</sub> is S or V, X<sub>4</sub> is D or E, X<sub>5</sub> is Q, K or R, and X<sub>6</sub> is L or V;

wherein said polypeptide further comprises a detectable label, and detecting cells which bind said labeled polypeptide.

31. The method of Claim 30 wherein said detectable label is a fluorescent label and said detecting is by fluorescence activated cell sorting.

5 32. A composition of matter comprising an Hsp47-related polypeptide pharmaceutically acceptable carrier.

33. An Hsp47 polypeptide comprising the sequence AVL<sub>1</sub>SAEQ<sub>1</sub>LR and a label.

10 34. The Hsp47 polypeptide of Claim 33 wherein said label is a fluorescent label.

35. An Hsp47-related immunoprotective polypeptide in combination with a cell, tissue or organ.

36. An antibody which specifically binds to an Hsp47 polypeptide epitope comprising the sequence AX<sub>1</sub>X<sub>2</sub>X<sub>3</sub>AEQ<sub>1</sub>LR.

15 37. An anti-idiotypic antibody to a polypeptide comprising the sequence AVL<sub>1</sub>SAEQ<sub>1</sub>LR.

38. A vector comprising nucleic acid encoding human Hsp47 polypeptide operably linked to control sequences recognized by a host cell transformed with the vector.

20 39. A host cell comprising the vector of Claim 38.

40. A process for producing an Hsp47 polypeptide comprising culturing the host cell of Claim 39 under conditions suitable for expression of said Hsp47 polypeptide and recovering said Hsp47 polypeptide from the cell culture.

5 41. A chimeric molecule comprising a heterologous amino acid sequence fused to an Hsp47 polypeptide.

42. The chimeric molecule of Claim 41, wherein said heterologous amino acid sequence is an Fc region of an immunoglobulin or biotin.

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## Signal Peptide

## Mature

M R S L L L L G T L C L L A V A L A | A E V K K  
 ATGCGCTCTCTCCTTCTGGGCACCTTATGCCTCCTGGCTGTGGCCCTGGCAGCCGAGGTGAAGAAA  
 TACGCGAGAGAGGAAGACCCGTGGAATACGGAGGACCGACACCGGGACCGTCCGCTCCACTTCTTT  
 P V E A A A P G T A E K L S S K A T T L A E  
 CCTGTAGAGGCCGAGCCCCCTGGTACTGCGGAGAAGCTGAGTTCCAAGGCGACCACTGGCAGAG  
 GGACATCTCCGGCGTCGGGGACCATGACGCCTCTTCGACTCAAGGTTCCGCTGGTGTGACCGTCTC  
 P S T G L A F S L Y Q A M A K D Q A V E N I  
 CCCAGCACAGGCCTGGCCTTCAGCCTGTATCAGGCAATGGCCAAGGACCAGGAGTGGAGAACATC  
 GGGTCGTGTCCGGACCGGAAGTCGGACATAGTCCGTTACCGGTTCCCTGGTCCGTCACCTCTTGTAG  
 L V S P V V V A S S L G L V S L G G K A T T  
 CTGGTGTACCCGTGGTGGTGGCCTCGTCGCTGGGTCTCGTGTGCTGGGCGGCAAGGCGACCACG  
 GACCACAGTGGGCACCACCACCGGAGCAGCGACCCAGAGCACAGCGACCCGCCGTTCCGCTGGTGC  
 A S Q A K A V L S A E Q L R D E E V H A G L  
 GCGTCGCAGGCCAAGGCAAGTGTGAGCGCCGAGCAGCTGCGCGACGAGGAGGTGCACGCCGGCCTG  
 CGCAGCGTCCGGTTCGCTCACGACTCGCGGCTCGTCGACGCGCTGCTCCTCCACGTGCGGCCGGAC  
 G E L L R S L S N S T A R N V T W K L G S R  
 GGTGAGCTGCTGCGCTCACTCAGCAACTCGACGGCGCGCAACGTGACCTGGAAGCTGGGCAGCCGA  
 CCACTCGAGCAGCGAGTGAGTCGTTGAGCTGCCGCGGCTTGCACTGGACCTTCGACCCGTCGGCT  
 L Y G P S S V S F A D D F V R S S K Q H Y N  
 CTGTACGGACCCAGCTCAGTGAGCTTTCGCTGATGACTTCGTGCGCAGCAGCAAGCAGCACTACAAC  
 GACATGCCTGGGTCGAGTCACTCGAAGCGACTACTGAAGCACGCGTCGTCGTTTCGTCGTGATGTTG  
 C E H S K I N F P D K R S A L Q S I N E W A  
 TGCGAGCACTCCAAGATCAACTTCCCGGACAAGCGCAGCGCGCTGCAGTCCATCAACGAGTGGGCC  
 ACGCTCGTGAGGTTCTAGTTGAAGGGCCTGTTTCGCGTCGCGCGACGTCAGGTAGTTGCTCACC CGG  
 A Q T T D G K L P E V T K D V E R T D G A L  
 GCGCAGACCACCGACGGCAAGCTGCCCGAGGTCACCAAGGACGTGGAGCGCACGGACGGCGCCCTG  
 CGCGTCTGGTGGCTGCCGTTTCGACGGGCTCCAGTGGTTTCCTGCACCTCGCGTGCCTGCCGCGGGAC  
 L V N A M F F K P H W D E K F H H K M V D N  
 CTAGTCAACGCCATGTTCTTCAAGCCACACTGGGATGAGAAATTCCACCACAAGATGGTGGACAAC  
 GATCAGTTGCGGTACAAGAAGTTCGGGTGTGACCCTACTCTTTAAGGTGGTGTCTTACCACCTGTTG  
 R G F M V T R S Y T V G V T M M H R T G L Y  
 CGTGGCTTCATGGTGA CTGCGTCTATACTGTGGGTGTTACGATGATGCACCGGACAGGCCTCTAC  
 GCACCGAAGTACCACTGAGCCAGGATATGACACCCACAATGCTACTACGTGGCCTGTCCGGAGATG  
 N Y Y D D E K E K L Q L V E M P L A H K L S  
 AACTACTACGACGACGAGAAGGAGAAGCTGCAGCTGGTGGAGATGCCCCTGGCTCACAAGCTCTCC  
 TTGATGATGCTGCTGCTCTTCTCTTCGACGTCGACCACCTCTACGGGGACCGAGTGTTCGAGAGG  
 S L I I L M P H H V E P L E R L E K L L T K  
 AGCCTCATCATCCTCATGCCCCATCACGTGGAGCCTCTCGAGCGCCTTGAAAAGCTGCTAACCAAA  
 TCGGAGTAGTAGGAGTACGGGGTAGTGACCTCGGAGAGCTCGCGGAACTTTTCGACGATTGGTTT  
 E Q L K I W M G K M Q K K A V A I S L P K G  
 GAGCAGCTGAAGATCTGGATGGGGAAGATGCAGAAGAAGCTGTTGCCATCTCCTTGCCCAAGGGT  
 CTCGTCGACTTCTAGACCTACCCCTTCTACGTCTTCTTCCGACAACGGTAGAGGAACGGGTTCCCA  
 V V E V T H D L Q K H L A G L G L T E A I D  
 GTGGTGGAGGTGACCCATGACCTGCAGAAACACCTGGCTGGGCTGGGCCTGACTGAGGCCATTGAC  
 CACCACCTCCACTGGGTACTGGACGTCTTTGTGGACCGACCCGACCCGGACTGACTCCGGTAAC TG  
 K N K A D L S R M S G K K D L Y L A S V F H  
 AAGAACAAGGCCGACTTATCACGCATGTCTGGCAAGAAGGATCTGTACCTGGCCAGTGTGTTCCAC  
 TTCTTGTTCCGGCTGAATAGTGC GTACAGACCGTTCTTCTTAGACATGGACCGGTCACACAAGGTG  
 A T A F E L D T D G N P F D Q D I Y G R E E  
 GCCACCGCCTTTGAGTTGGACACAGATGGCAACCCCTTTGACCAGGACATCTACGGGCGCGAGGAG  
 CGGTGGCGGAACTCAACCTGTGTCTACCGTTGGGGAACTGGTCTGTAGATGCCCGCGCTCCTC  
 L R S P K L F Y A D H P F I F L V R D T Q S  
 CTGCGCAGCCCCAAGCTGTTCTACGCCGACCACCCCTTCATCTTCTGGTGGCGGGACCCCAAGC  
 GACGCGTCGGGGTTCGACAAGATGCGGCTGGTGGGGAAGTAGAAGGACCACGCCCTGTGGGTTTCG  
 G S L L F I G R L V R L K G D K M R D E L .  
 GGCTCCCTGCTATTTCATTGGGCGCCTGGTCCGGCTCAAGGGTGACAAGATGCGAGACGAGTTATAG  
 CCGAGGGACGATAAGTAACCCGCGGACCAGGCCGAGTTCCCACTGTTCTACGCTCTGCTCAATATC

**FIG. 1**

SUBSTITUTE SHEET (RULE 26)



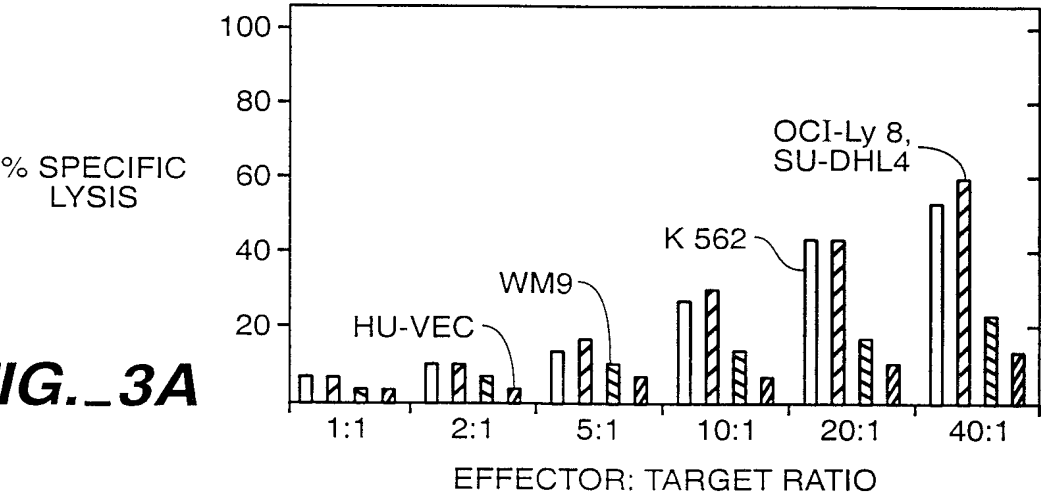
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Alignment of HSO47 with HLA/IL-12 $\beta$  Chains

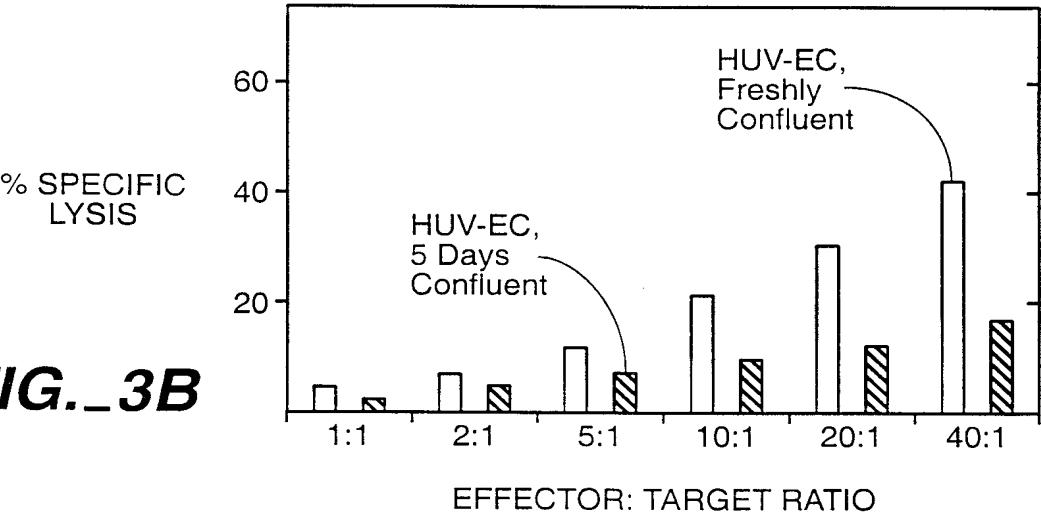
<u>Molecule</u>	<u>Position</u>	<u>Amino Acid Sequence</u>	<u>Position</u>
Human Hsp47	96	AvlsAEQLR	104
Hu Colligin 2	95	AVLSAEQLR	103
HLA-A2 A*0201	149	AahvAEQLR	157
HLA-A2 A*0204	167	AahvAEQLR	175
HLA-A2 A*0206	175	AahvAEQLR	183
HLA-A2 A*0211	166	AahvAEQLR	174
HLA-A10 alpha	84	AahvAEQLR	92
HLA-Aw 69	150	AahvAEQLR	158
Rat Hsp47	94	AVLSAEkLR	102
Mouse Hsp47	94	AVLSAEkLR	102
Hamster Hsp47	94	AVLSAEkLR	102
Chicken Hsp47	82	AVLSAdkLn	90
Hu IL-12 $\beta$ -chain	150	AtLSAErvR	158
mouseIL1 2 $\beta$ -chain	150	AtLSAErvR	158
cat IL-12 $\beta$ -chain	150	AtLSAEkvR	158
cow IL12 $\beta$ -chain	150	AILSAEkvn	158
Consensus HSP		AVLSA(d,e)(k,q)LR	
Consensus HLA		A(v,a)(l,h)(a,v)AEQLR	
Consensus IL-12 $\beta$ -chain		A(v,l,t)LSAE(q,k,r)(l,v)R	
Overall Consensus		A(v,l,a,t)(l,h)(a,v)A(d,e)(k,q,r)(l,v)R	

**FIG.\_2**

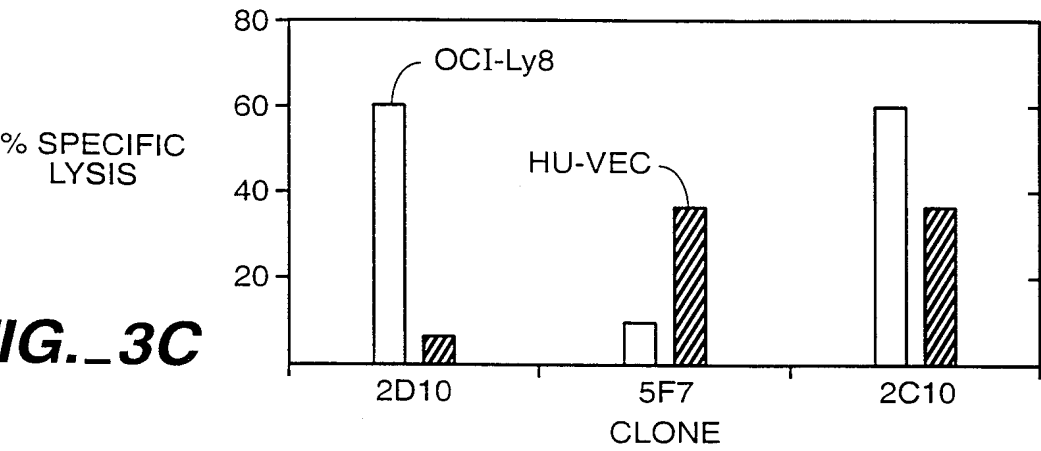
**FIG.\_3A**



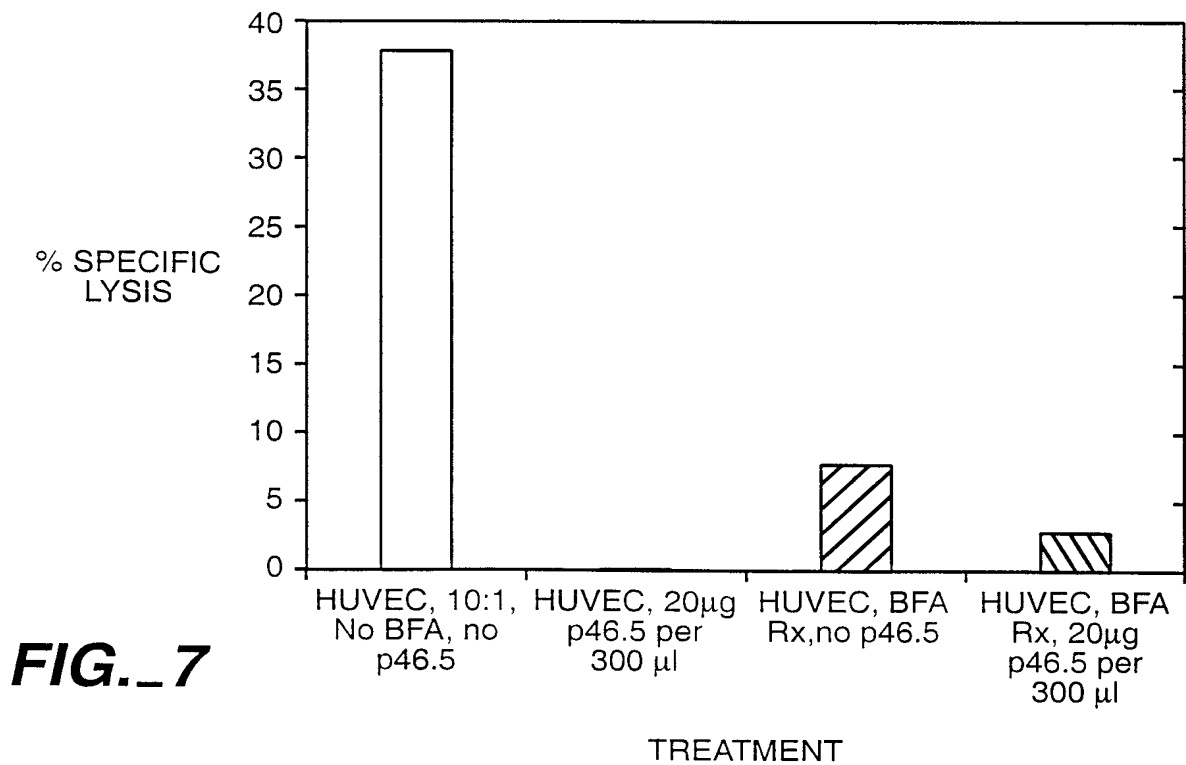
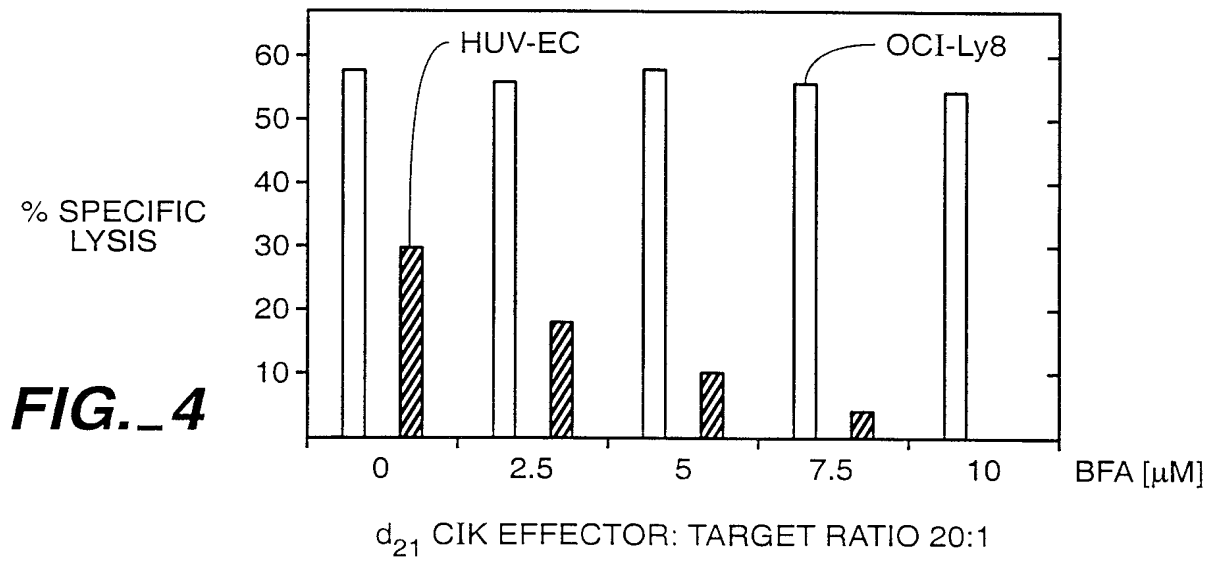
**FIG.\_3B**



**FIG.\_3C**



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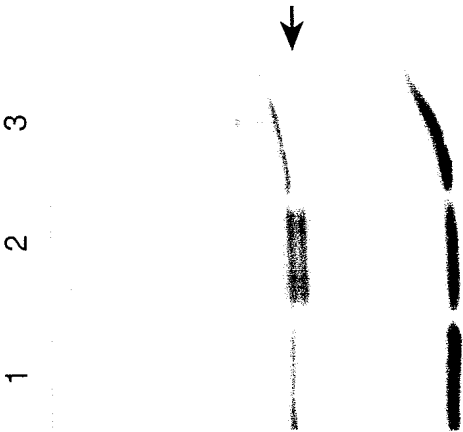


FIG.\_5C

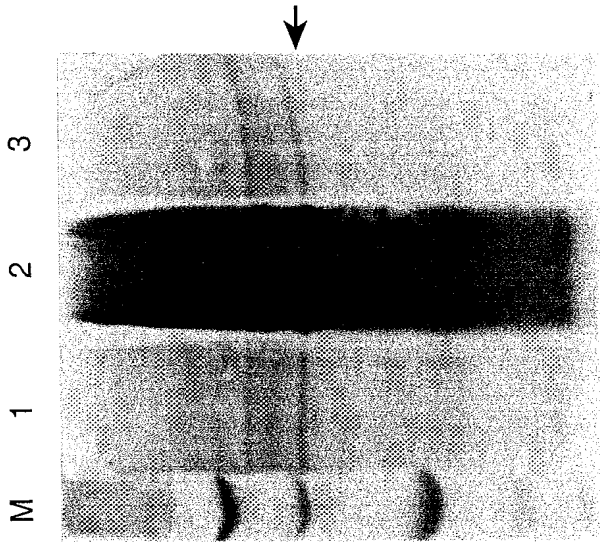


FIG.\_5B

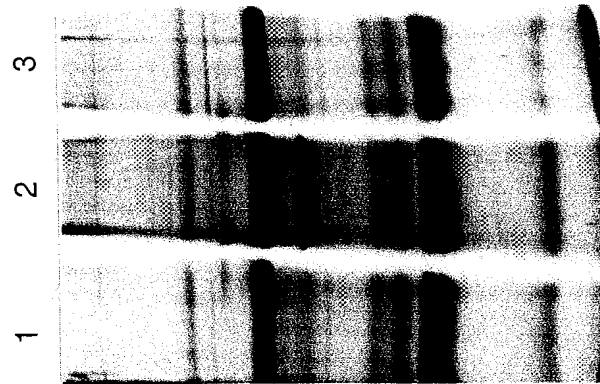
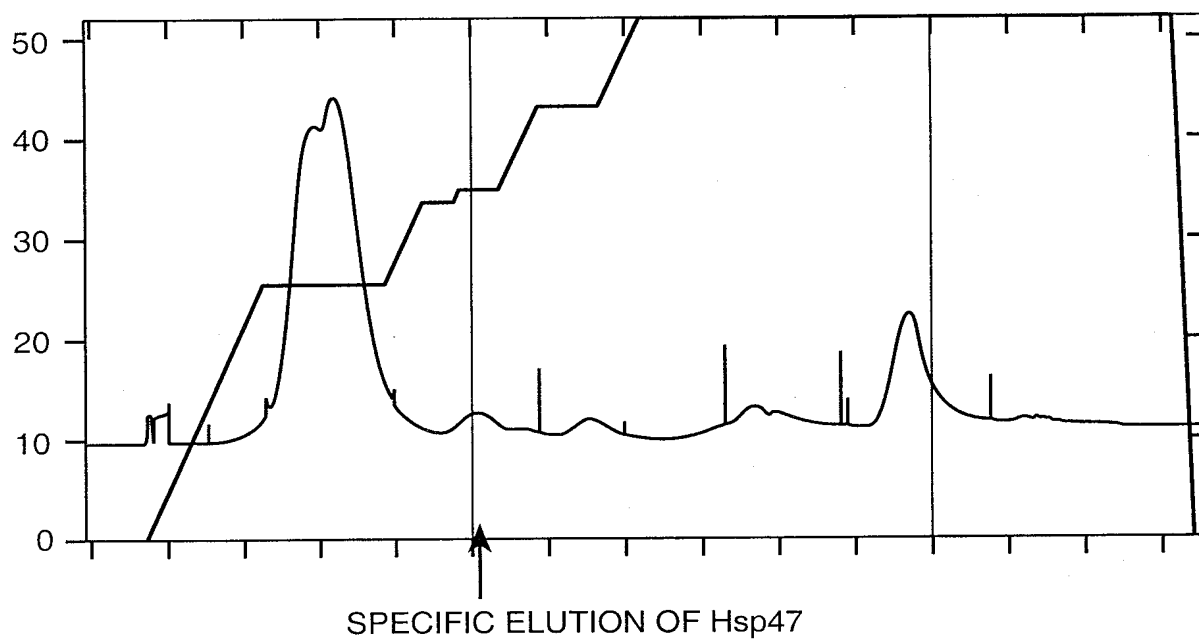
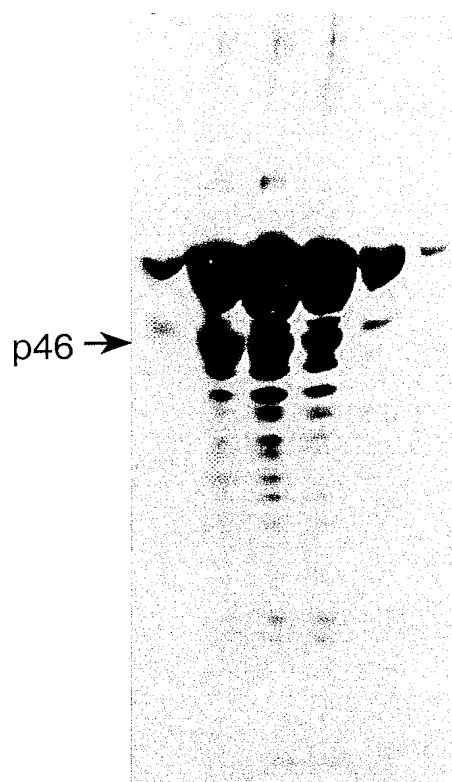
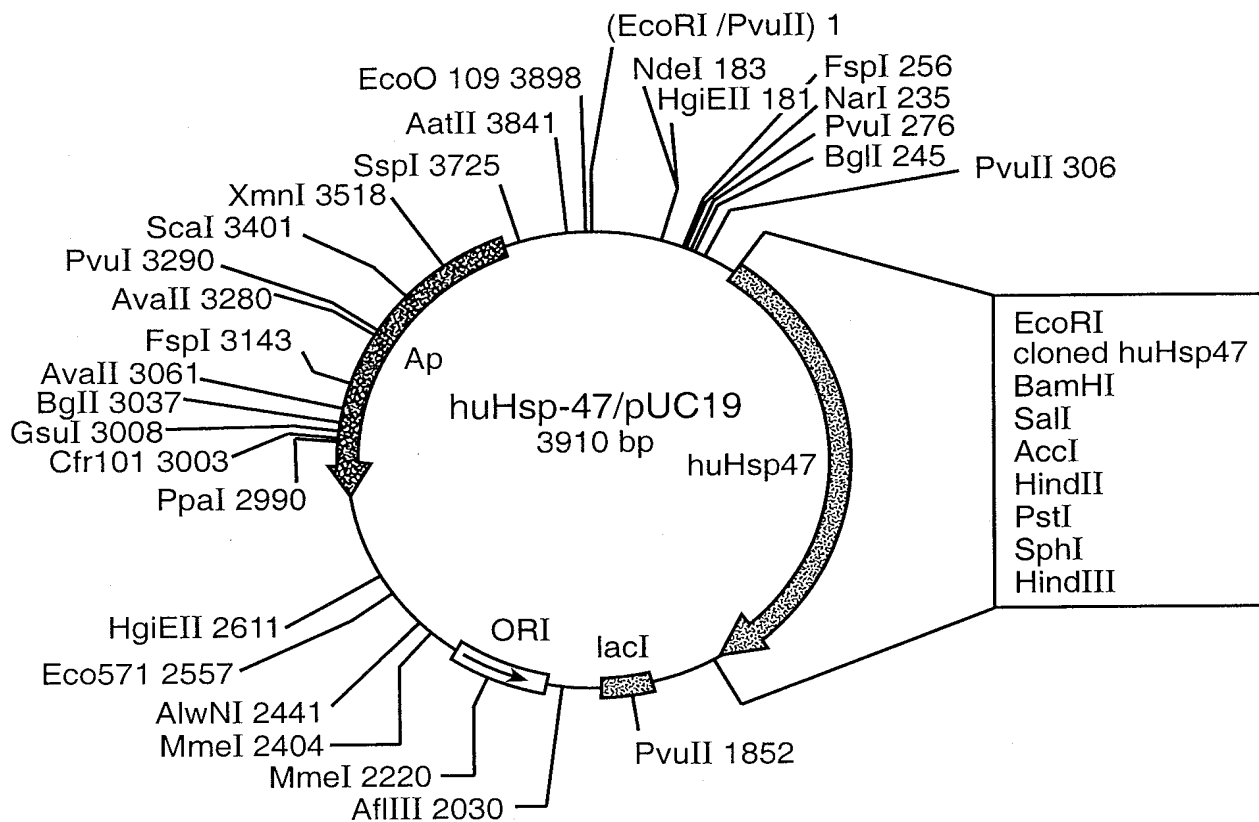
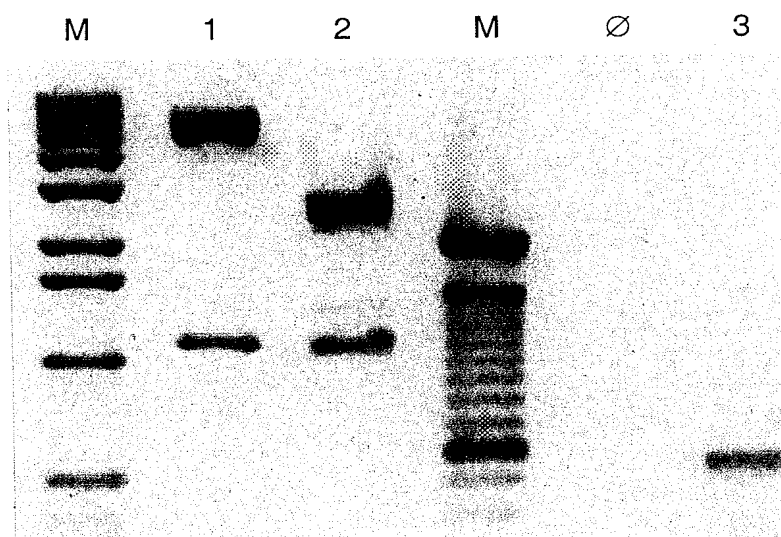


FIG.\_5A

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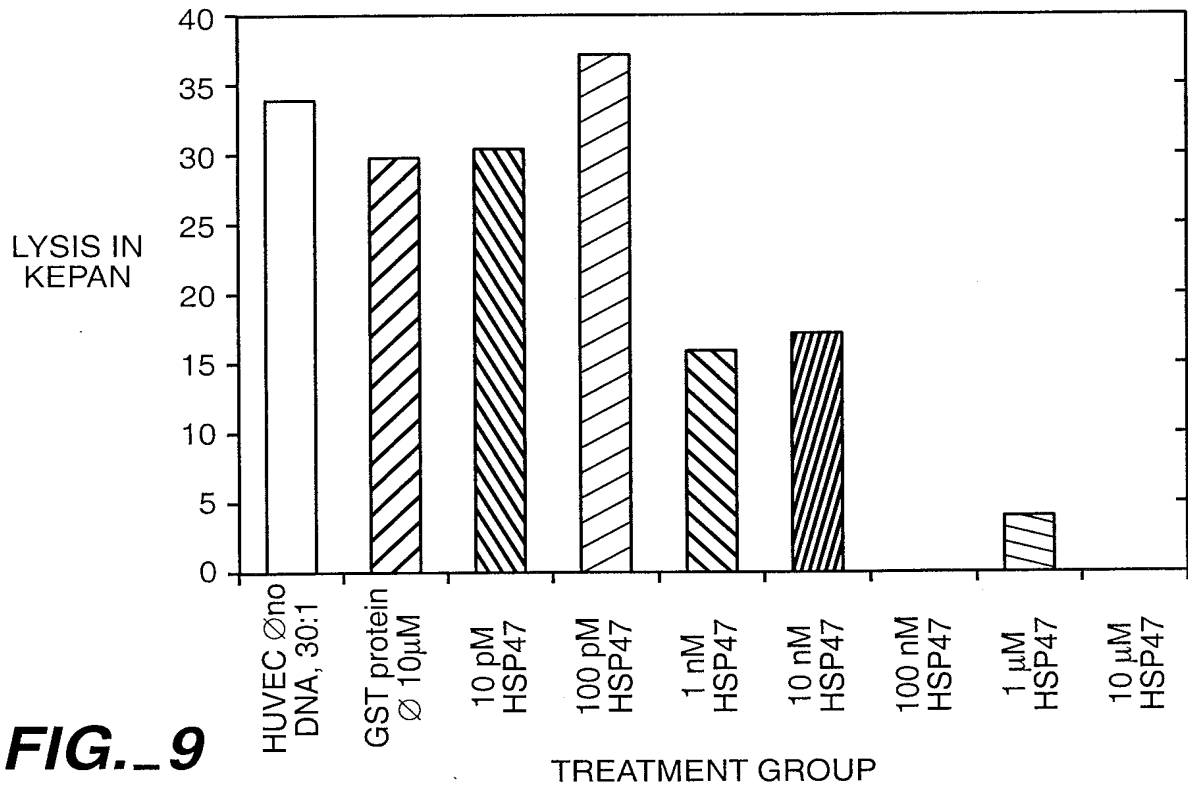
**FIG.\_6A****FIG.\_6B**

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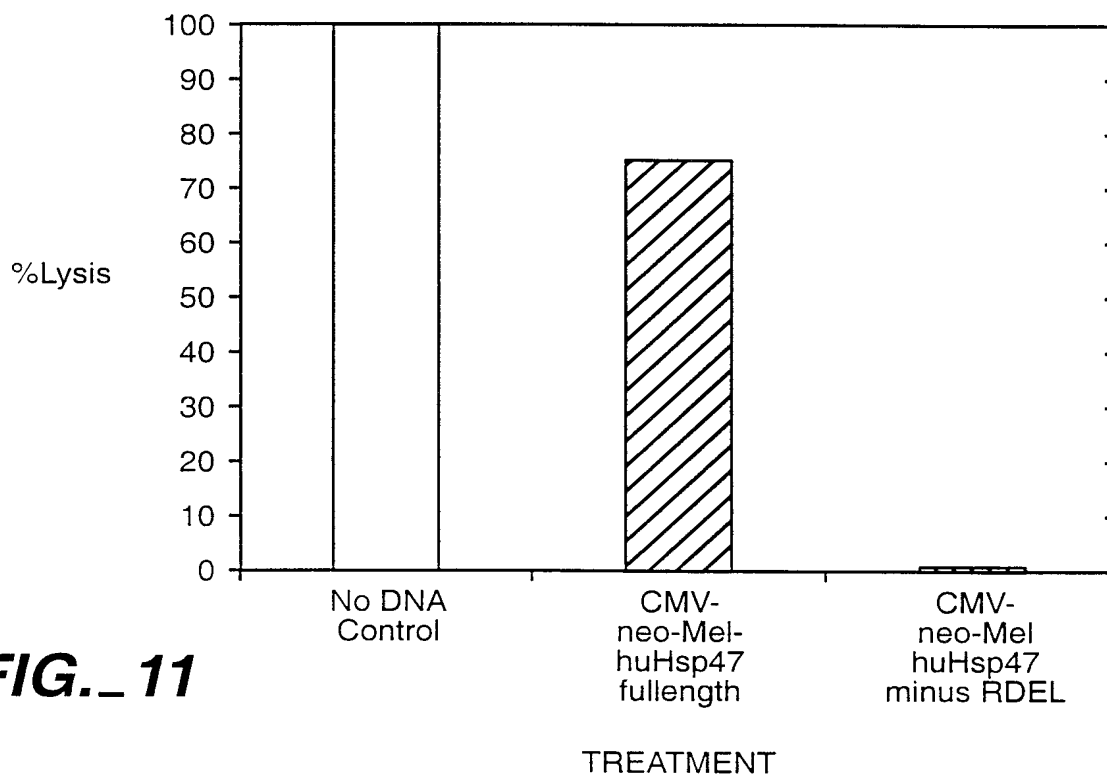
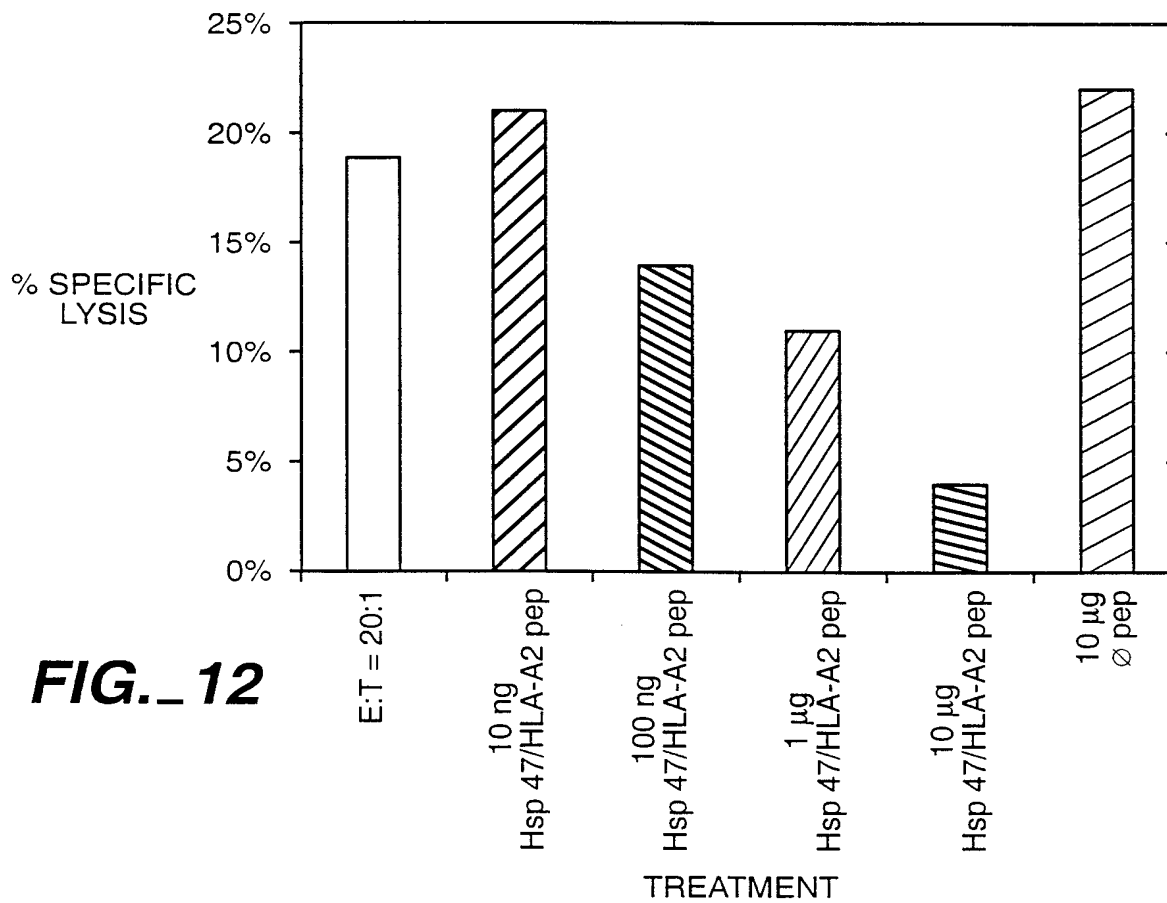
**FIG.\_8A****Plasmid name:** huHsp47/pUC19**Plasmid size:** 3910 bp**Constructed by:** Ernest-G. Hope**Construction date:** 1996

**Comments/References:** puc 19 based huHsp47 gene cassette derived from 2 partial clones in pCR1000/pGEM via primers (A. 5' -ACGTTTGGATCCAGGTGAAGA, 3' -GTCCTTGGCCAT B. 5' GCAATGGCCAAGGACCAGGCAGTGGAG, 3' ATCTGAATTCCTATAACTCGTCTCGCA)

**FIG.\_8B**



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**FIG. 11****FIG. 12**



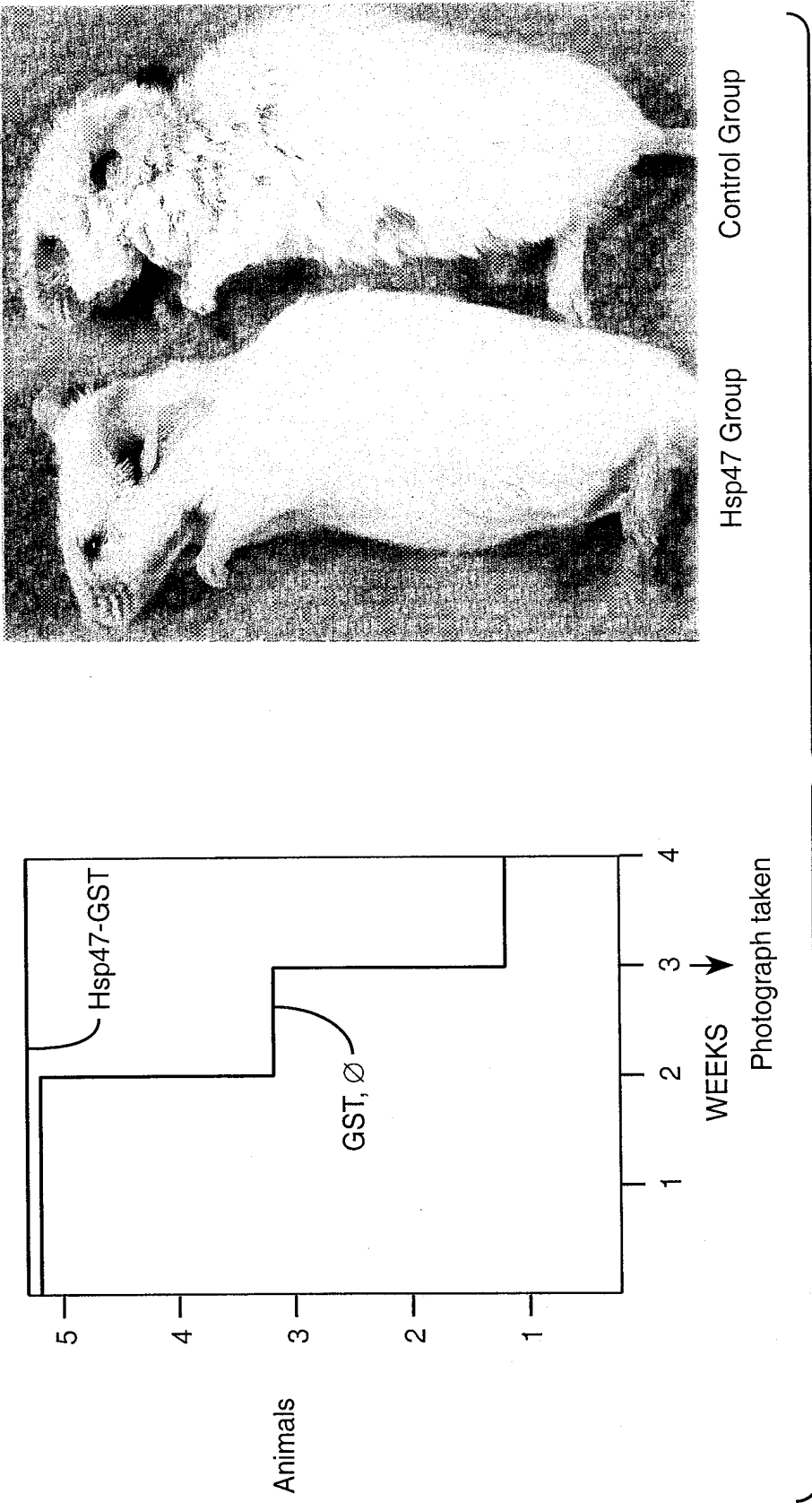
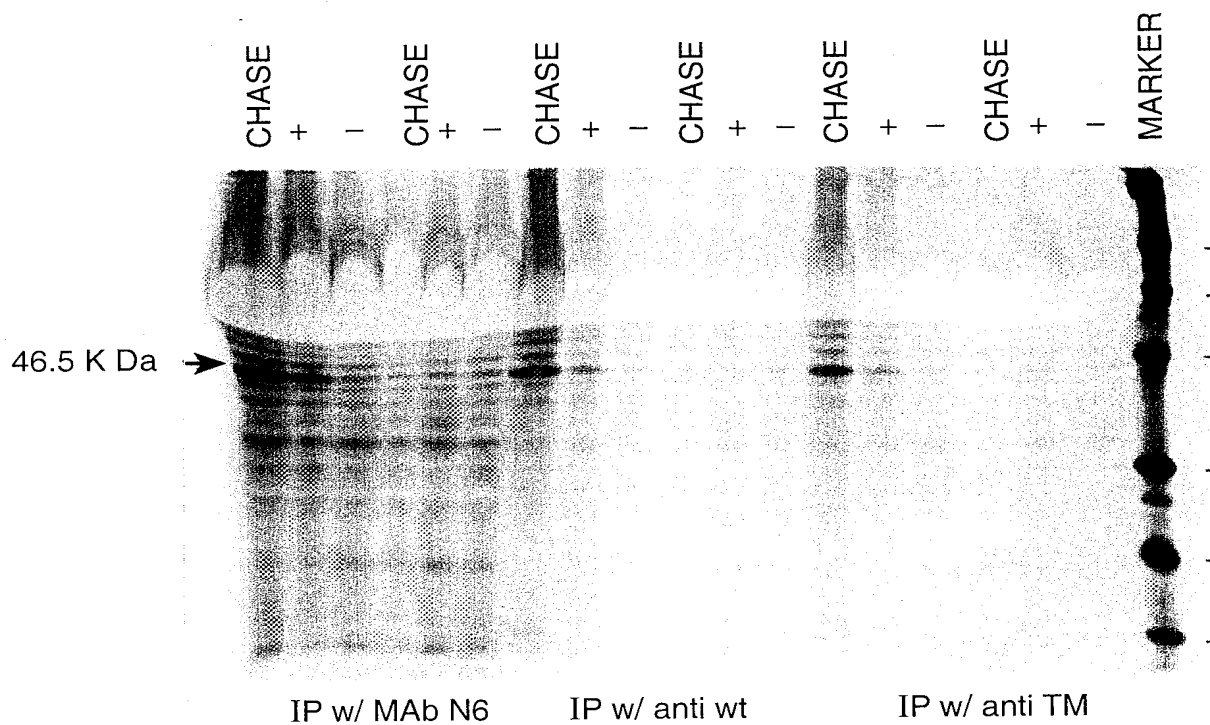
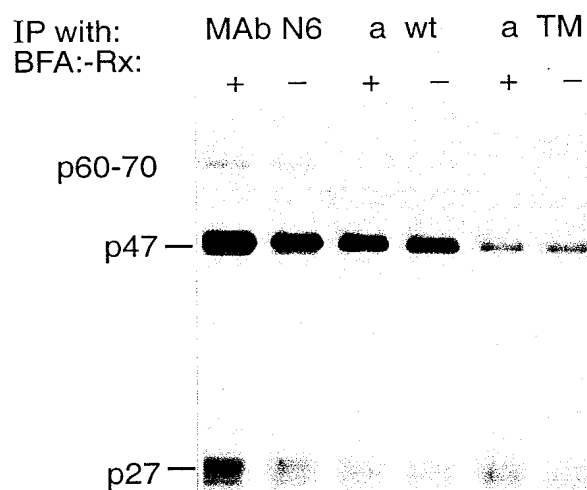


FIG. 13

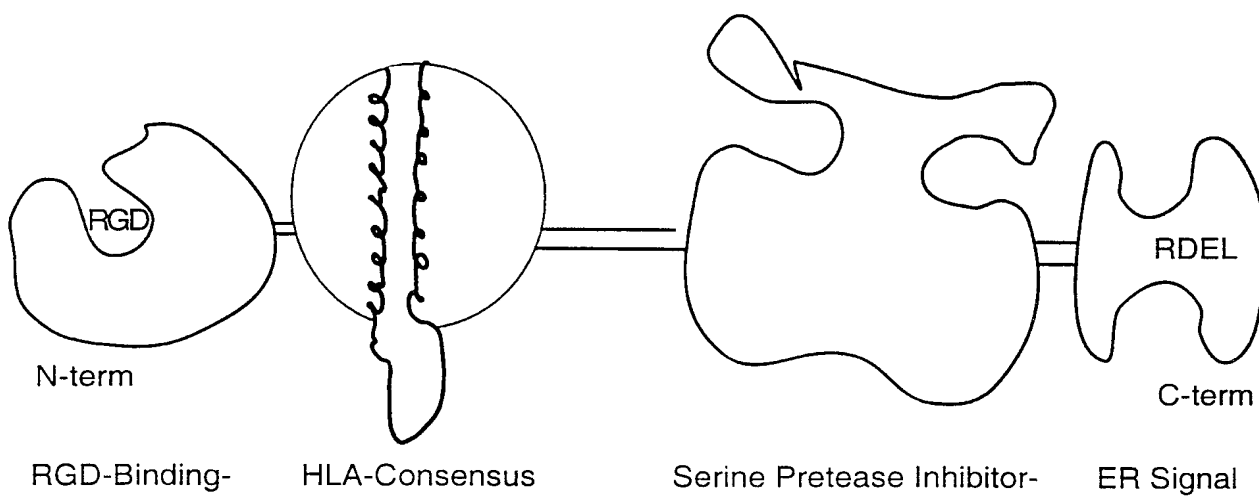
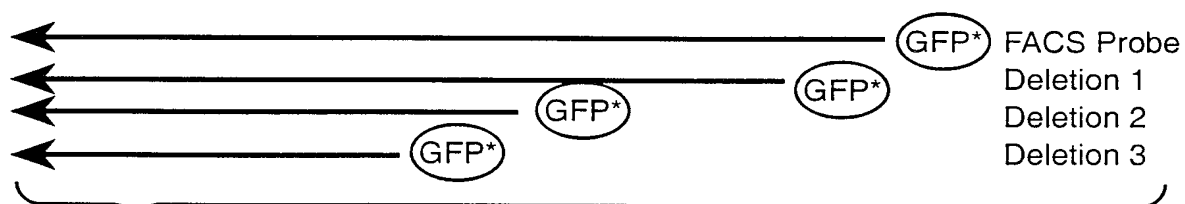
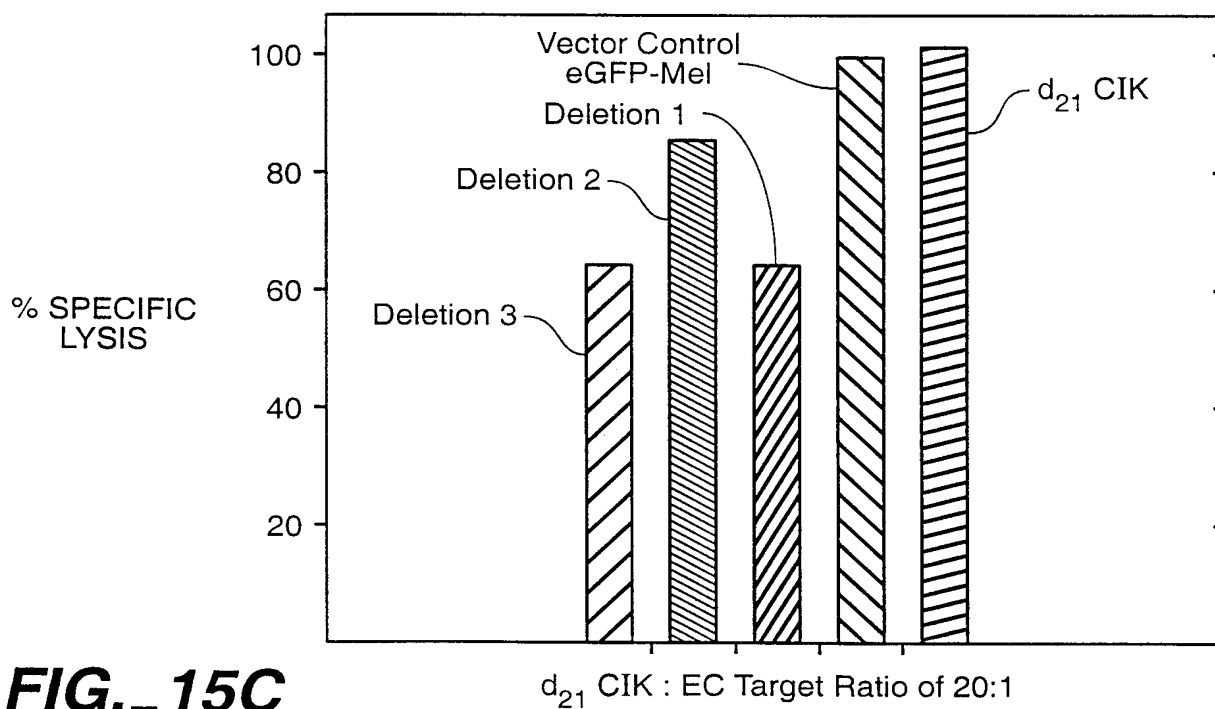


**FIG.\_14A**



**FIG.\_14B**

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**FIG.\_ 15A****FIG.\_ 15B****FIG.\_ 15C**

## INTERNATIONAL SEARCH REPORT

International application No.  
PCT/US99/19337

**A. CLASSIFICATION OF SUBJECT MATTER**

IPC(7) : A61K 39/395, 38/02; C07K 16/18; C12N 1/21, 15/85; C12Q 1/68

US CL : Please See Extra Sheet.

According to International Patent Classification (IPC) or to both national classification and IPC

**B. FIELDS SEARCHED**

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 514/2, 530/387.3, 435/6, 320, 252.3, 325, 424/192.1; A61K 39/395

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

STNCAS patents medline embase biosis caplu

**C. DOCUMENTS CONSIDERED TO BE RELEVANT**

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	SALOH et al. Intracellular Interaction of Collagen Specific Stress Protein HSP47 with Newly Synthesized Procollagen. The Journal of Cell Biology. 1996, Vol. 133, No. 2, pages 469-483, especially pages pgs 470-471.	32, 35, 38-42
Y	VERRICO et al. Expression of the Collagen-related heat shock protein HSP47 in fibroblasts treated with hyperthermia or photodynamic therapy. British Journal of Cancer. 1997, Vol. 76, No. 6, pages 719-724, especially page 720.	38-40
Y	EP 0 813 871 A1 (KUREHA CHEMICAL INDUSTRY, CO.) 12 December 1997, page 8, lines 22-25.	32 and 35

☐ Further documents are listed in the continuation of Box C. ☐ See patent family annex.

* Special categories of cited documents:	*T* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
*A* document defining the general state of the art which is not considered to be of particular relevance	*X* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
*E* earlier document published on or after the international filing date	*Y* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
*L* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	*G* document member of the same patent family
*O* document referring to an oral disclosure, use, exhibition or other means	
*P* document published prior to the international filing date but later than the priority date claimed	

Date of the actual completion of the international search

17 JANUARY 2000

Date of mailing of the international search report

10 FEB 2000

Name and mailing address of the ISA/US  
Commissioner of Patents and Trademarks  
Box PCT  
Washington, D.C. 20231

Facsimile No. (703) 305-3230

Authorized officer

AMY DECLoux

Telephone No. (703) 308-0196

# INTERNATIONAL SEARCH REPORT

International application No.  
PCT/US99/19337

## Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:  
because they relate to subject matter not required to be searched by this Authority, namely:
  
2. ☒ Claims Nos.: 2-5, 9-13, 20-23, 27-31, 33-34, 36-37  
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:  
  
No CRF for this case has been filed. The instant claims recite sequences or depend therefrom and cannot be searched other than by a sequence search.
  
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
  
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
  
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
  
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.  
☐ No protest accompanied the payment of additional search fees.

# INTERNATIONAL SEARCH REPORT

International application No.

PCT/US99/19337

## A. CLASSIFICATION OF SUBJECT MATTER:

US CL :

514/2, 530/387.3, 435/6, 320, 252.3, 325, 424/192.1; A61K 39/395